

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 118984

TO: Sean McGarry

Location: REM-2D/19/2C18

Art Unit: 1635

Wednesday, April 07, 2004

Case Serial Number: 09/993731

From: Paul Schulwitz

Location: Biotech-Chem Library

REM-1A65

Phone: (571)272-2527

paul.schulwitz@uspto.gov

Search Notes

Examiner McGarry,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Paul Schulwitz Technical Information Specialist STIC Biotech/Chem Library (571)272-2527



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21 bp from Patent WO0138576.

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PAT 22-JUN-2001

Page 2

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ALIGNMENTS

SOURCE ORGANISM Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; 21 bp DNA 6 from Patent WO0138576. 0.8%; Score 20.6; DB 1; 95.2%; Pred. No. 6.3; vative 1; Mismatches 0, rganism="Homo sapiens" >1_type="unassigned DNA" >_xref="taxon:9606" Ireland, J.S. and Lander, E.S.
e nucleotide polymorphisms
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cation/Qualifiers 0.8%; Score 20.6; DB 1; 95.2%; Pred. No. 6.3; vative 1; Mismatches 0; Ireland, J.S. and Lander, E.S.

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1138576-A 155 31-MAY-2001;
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cation/Qualifiers etazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Ltheria; Primates; Catarrhini; Hominidae; Homo. RIGGAGCICIC 21 GTGGAGCTCTC 1843 rganism="Homo sapiens" ol_type="unassigned DNA" o_xref="taxon:9606" Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mtheria; Primates; Catarrhini; Hominidae; Homo. GI:14535671 GI:14535672 TCAGCAGGCCC 1644 rcadcaddccc 21 21 bp from Patent W00138576. (human) DNA 0 Length 21; Length 21; Indels Indels linear linear PAT 22-JUN-2001 PAT 22-JUN-2001 0 0 Gaps Gaps 0, 0

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Human single nucleotide polymorphisms
Patent: WO 0138576-A 156 31-MAY-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
Location/Qualifiers
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Crouzet,J., Scherman,D. and Wils,P.
PURIFICATION OF A TRIPLE HELIX FORMATION WITH AN IMMOBILIZED OLIGONUCLECTIDE
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Other publication AU 4178996 960703
Other publication FR 2728264 960621.
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Patent: US 6319672-A 18 20-NOV-2001;
Location/Qualifiers
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/mol_type="unassigned DNA"
                                                                                                                                           Location/Qualifiers
                                                                                              /organism="synthetic construct"
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Sequence 18 from Patent EP1281774.
AX686853
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Matson,R.S., Coassin,P.J., Rampal,J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 41 09-NOV-1999;
Location/Qualifiers
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Patent: EP 1281774-A 18 05-FEB-2003;
Aventis Pharma S.A. (FR)
1 (bases 1 to 21)
Matson,R.S., Coassin,P.J., Rampal,J.B. and Oligonucleotide repeat arrays
Patent: US 5981185-A 53 09-NOV-1999;
Location/Qualifiers
1. .21
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                           organism="unknown"
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                                              Caskey, C. Thomas.
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Sequence 59
AR084570
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Matson,R.S., Coassin,P.J., Rampal,J.B. and
Oligonucleotide repeat arrays
Patent: US 5981185-A 59 09-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unknown
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Matson,R.S., Coassin,P.J., Rampal,J.
Oligonucloctide repeat arrays
Patent: US 5981185-A 64 09-NOV-1999;
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/mol_type="unassigned DNA"
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/mol_type="unassigned
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/mol_type="unassigned DNA"
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from patent US 5756684.
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AR034773
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unclassified.
1 (bases 1 to 24)
Johnson, E.M. and Bergemann, A.D.
                                                                    Lipford, G.B., Heeg, K. and Wagner, H. G-motlif oligomuclectides and uses thereof Patent: WO 0014217-A 39 16-MAR-2000; LIPFORD GRAYSON B (DE) ; HEEG KLAUS (DE) CPG IMMUNOPHARMACEUTICALS GMBH (DE)
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AX023424
AX023424.1 GI:10183824
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/noCe="synthetic, no natural or
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Matson,R.S., Cosssin,P.J., Rampal,J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 83 09-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 21)
Matson,R.S., Coassin,P.J., Rampal,J.B.
Oligonuclectide repeat arrays
Patent: US 5981185-A 70 09-NOV-1999;
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/mol_type="unassigned DNA"
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/mol_type="unassigned
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                                                                 Roemer,T., Jiang,B., Boone,C. and Bussey,H. Gene disruption methodologies for drug target Patent: WO 0160975-A 397 23-AUG-2001; Elitra Pharmaceuticals, Inc. (US)
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Dean,N. and Cowsert,L.M.
Antisense modulation of daxx expression
Patent: US 6180353-A 153 30-JAN-2001;
Location/Qualifiers
                                                                                                                                                                              Sequence 397 from Patent AX241159
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Sequence 153 from
AR126724
                                                                                                                                      synthetic construct synthetic construct
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/mol_type="unassigned DNA"
                            organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                    'note="DNA primer"
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18; Conserv
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Candida albicans
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AX154328
AX154328.1 GI:14535942
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Patent: WO 0138576-A 426 31-MAY 2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RE
                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo-
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/mol_type="unassigned DNA"
/db_xref="taxon:5476"
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                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="unassigned DN
/db_xref="taxon:9606"
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Sequence 12 from patent US 6159710.
AR121232
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Ward, D.T. and Cowsert, L.M.
Antisense modulation of NCK-2 expres
Patent: US 6165728-A 54 26-DEC-2000;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 21)
Fraser.N.W., Zabolotny,J.M. and Krummenacher,C.F.
Method and compositions for stabilizing unstable gene transcripts
Patent: US 6159710-A 12 12-DBC-2000;
Location/Qualifiers
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                                              Varshney, G.C.
Specific mucin expression as a marker
Patent: US 6576423-A 22 10-JUN-2003;
                                                                          Unclassified.

1 (bases 1 to 18)

Batra, S.K., Brand, R.E., Ringel, J., Faulmann, G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                   Unclassified
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/mol_type="unassigned DNA"
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/mol_type="unassigned
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/mol_type="genomic
            organism="unknown"
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al Similarity 94.4%;
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1 (bases 1 to 20)

MCKay,R., Freier,S.M. and Wyatt,J.

Antisense modulation of casein kinase 2-alpha

Antisense modulation of 24-SEP-2002;

Patent: US 6455307-A 93 24-SEP-2002;

Location/Qualifiers
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17; Conserv
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Sequence 173 from patent
AR307962
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Dean,N.M. and Cowsert,L.M.
Antisense modulation of daxx expression
Patent: US 6180353-A 155 30-JAN-2001;
Location/Qualifiers
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/mol_type="genomic DNA"
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                                    CE 1 (bases 1 to 17)

RS Blatt, L. Zwick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic acid molecules

Patent: UP 2002541795-A/2806

SELWARYOTE PA/2806

PO 10-DEC-2002

PF 11-APR-2000 UP 2000611654

PF 11-APR-2000 UP 2000611654

PF 11-APR-1999 US 60/129390

PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN PC C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC C12P21/02,
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Regulation of repressor genes
BD255013
BD255013.1_GI:33064783
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Sequence 174 from patent
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PC
C12P21/02,C12P21/02//A61K31/711,(C12N5/10,C12R1:91),(C12P21/02,
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/mol_type="genomic |
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/mol_type="genomic |
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126890/c
LOCUS
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I91631/c
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191631.1 GI:3
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126890
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Sidransky,D.
Nucleic acid mutation detection by analysis of Patent: US 5561041-A 113 01-OCT-1996;
Location/Qualifiers
1. .17
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FT Source 1..17
FT Source /organism='Eukaryote'.
                                                                            Patent:
                                                          Analysis of sputum by amplification an acid sequences
Patent: US 5726019-A 113 10-MAR-1998;
Location/Qualifiers
1. 17
                                                                                                                1 (bases 1 to 17)
Sidransky, D.
                                                                                                                                                    Unknown
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/mol_type="genomic DNA"
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                                                                                                                                                                                       GI:3936101
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Score 15.4; DB Pred. No. 32;
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Patent: WO 0159103-A 2359 16-AUG-2001;
RIBOZYME PARAMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2359 from Patent AX216917
                                                    16;
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Sequence 567 from Patent W00188124.
AX422231
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                                                                                                                                                                  Method and reagent for the inhibition of erg
Patent: WO 0188124-A 567 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chórdata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                       Jarvis,T., von Carlowitz,T., Mcswiggen,J.A., Mclaughlin,F.G. Randi,A.M.
                                                                                                                                                                                                                                                                      Homo sapiens
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Similarity 94.1%;
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                                                                                                 /organism="Homo sapiens"
/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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/mol_type="unassigned RNA"
/db_xref="taxon:32630"
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PC C07K1
PC C07K1
PC 02,C1
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16; Conserv
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AX498864
AX498864.1 GI:23381157
 1 Similarity
                                                                                                                                                                                                                                                                 OS Homo sapiens (human)

PN 102526100-A/20

PN 202526100-A/20

PD 20-AUG-2002

FR 07-OCT-1999 JP 2000574561

FR 07-CCT-1999 US 60/103322,30-OCT-1998 US 60/106473 |

PR 07-DCC-1998 US 66/114088

PI MARGARET ANN JCHNS, BRIAN MOLDOVER, JAMES DAVID OFFORD PC C12N15/09, A61K31/711, A61K38/00, A61K48/00, A61P25/06, A61P25/08,
                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 19) Johns, M.A., Moldover, B. and Offord, J.D.
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JP 2002526100-A/20.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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larity 94.1%; Pred. No. 32;
Conservative 0; Mismatches
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C07K14/47,
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                                                                                                          /organism='Homo
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                                                   /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 44;
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RESULT 38
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AUTHORS
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                                                                                                                                                                                                                                                                                                                        Sequence 46
AR078620
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Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.
Regulation of repressor genes using nucleic ac
Patent: JP 2002541195-A 2805 10-DEC-2002;
RIBOZYME PHARMACEUTICALS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unidentified
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                                                      Antisense modulation of RhoB expression Patent: US 5962672-A 46 05-OCT-1999; Location/Qualifiers 1..18
                                                                                                                                                        1 (bases 1 to 18)
Coswert, L.M.
                                                                                                                                                                                                                                        Unknown
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12-APR-1999 US 60/12990
LAWRENCE BLATT,MCHABEL ZWICK,PAMELA PAVCO,JAMES MCSWIGGEN
15/09,A61K38/00,A61K48/00,A61P43/00,A61P43/00,C12N5/10, PC
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JP 2002541795-A/2805
10-DEC-2002
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[] / Organism='Eukaryote'
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
/organism="unknown"
/mol_type="unassigned DNA"
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100.0%; Pred. No.
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RESULT 40
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AR215559 Sequence AR215559

107

18 bp I from patent US 6410323.

PAT

25-SEP-2002

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                       1821 GAGGCCGGCGAGGTGGAG 1838
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ANTIBIOTICOS
OS Tritirach
PN JP 200255
PD 10-DEC-20
PD 07-APR-11
PI JAVIER VI
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PR 09-APR-1999 ES P 9900731
PI JAVIER VELASCO ALVAREZ, SANTIAGO GUTIERREZ MARTIN, FRANCISCO JAVIER CASQUEIRO BLANCO, SONIA CAMPOY GARCIA, FRANCISCO FIERRO FIERRO, JOSE LUIS BARREDO FUENTE, BRUNO FRANCISCO FIERRO FIERRO, JOSE LUIS BARREDO FUENTE, BRUNO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tritirachium album
Tritirachium album
Eukaryota; Fungi; Ascomycota; mitosporic Ascomycota; Tritirachium.
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JP 2002541812-A/2.
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 GAGGCCGGCGACGTGGGG 18
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C12N15/09,C12N1/15,C12N1/19,C12N1/21,C12P35/06//(C12P35/06,
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JP 2002541812-A/2
                                                                                                                                                                                                                                sequence
(Pro-His-Val-Ala-Gly-Leu)
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                                                                                               /organism="Tritirachium
/mol_type="genomic DNA"
/db_xref="taxon:5558"
                                                                                                                                                  /organism='Tritirachium
Location/Qualifiers
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1363 CTGAGGCTTACCAGAAGC 1380
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16; Conserv
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1 (bases 1 to 18)

Roberts M.L. and Cowsert, L.M.

Antisense modulation of human Rho family gene expression
Patent: US 6410323-A 107 25-JUN-2002;
                                                                                                                                                                                                                                                                                                            Submitted (28-APR-1995) Yoshiyuki Sakaki, Institute of Medical Science, University of Tokyo, Human Genome Center; 4-6-1 Shirokanedai Minato-ku, Tokyo 108, Japan (E-mail:sakaki@igo.ims.u-tokyo.ac.jp, Tel:03-5449-5362, Fax:03-5449-546)
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sixty new STSs (sequence-tagged sites) of human chromosome DNA Res. 1 (2), 85-89 (1994) 96051984
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STS.
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                                                                                                                                                                                                                                                                                                                                                                                             2 (bases 1 to 18)
Sakaki,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 18)

Tanahashi, H., Ito, T., Hattori, M., Ohira, M., Ohki, M., Tashiro, K. as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                               Phone: 03-5449-5362
Fax : 03-5449-5445.
Location/Qualifiers
                                                                                                                                                                                                                                               Human Genome Center
Institute of Medical Science
University of Tokyo
                                                                                                                                                                                                                                                                                      Submitted (28-Apr-1995) to DDBJ by: Yoshiyuki Sakaki
                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
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/mol_type="genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                         /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                 chromosome="21"
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15; Conserv
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PR 27-SEP-2000 JP 00P 293692,06-FEB-2001 JP 01P
TORU TSUJI,HIROSHI YANAGAWA
PC C12N15/09,C12P21/02
CC Description of Artificial Sequence:Synthesized
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Method of constructing mutation DNA library and util:
Patent: WO 0226964-A 39 04-APR-2002;
MITSUBISHI CHEMICAL CORP.TORU TSUJI, HIROSHI YANAGAWA
OS Artificial Sequence
PN WO 0226964-A/39
PD 04-APR-2002
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synthetic construct
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Odink, K.G., Tarcsay, L.,
Sorg, C., DeWolf-Peeters,
                                                                                                                                                                                                                                                       synthetic construct
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artificial sequences.
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A34251
Synthetic sequencing primer.
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1 (bases 1 to 16)
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CIBA-GEIGY AG
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Of constructing mutation DNA library and utilization of constructing mutation DNA library and utilization
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism='Artificial
                                                                                               /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                             Location/Qualifiers
1. .17
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                                                         Score 14.4;
Pred. No. 49;
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AR189954
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AR286297/c
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AR189954
                                                                              Sequence 66
AR286297
AR286297.1
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 5442 12-FEB-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                               1 (bases 1 to 17)
Ward,D.T. and Cowsert,L.M.
Antisense modulation of RIP2 expression
Patent: US 6426221-A 4 30-JUL-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4 from patent US AR221454
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1 (bases 1 to 17)
Beigelman,L., Burgin,A., Beaudry,A., Karpeisky,A.,
Matulic-Adamic,J., Sweedler,D. and Zinnen,S.
                                  Unknown.
Unclassified.
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/mol_type="genomic
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RESULT 47
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1 Similarity 93.8%; Pred. No. 49;
15; Conservative
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Sequence 2336 from patent
AR324934
                                                                                                                               1 (bases 1 to 17)
Beigelman,L., Burgin,A.B., Beaudry,A., Karpeisky,A.,
Matulic-Adamic,J., Sweedler,D. and Zinnen,S.
Oligoribonucleotides with enzymatic activity
Patent: US 6617438-A 668 09-SEP-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 17)
Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 2336 20-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic ribonucleic acids with RNAse Patent: US 6528640-A 669 04-MAR-2003;
                                                15;
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Similarity 93.8%;
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/mol_type="unassigned RNA"
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                                                                                           /mol_type="unassigned RNA"
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AX216918
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AX216922
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17 bp
Sequence 915 from Patent W00173002.
AX263524
AX263524.1 GI:16512323
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Patent: WO 0159103-A 2364 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
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larity 93.8%; Pred. No. 49;
Conservative 0; Mismatches
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/db_xref="taxon:32630"
/note="Nucleic Acid"
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/mol_type="unassigned RNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                               /note="Nucleic Acid"
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Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae,
                                                   Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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AX263532
AX263532.1 GI:16512331
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Location/Qualifiers
Kmiec,E.B., Gamper,H.B. and Rice,M.C. Targeted chromosomal genomic alterations with modified single stranded oligonucleotides
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. "
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/mol_type="unassigned DNA"
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Sequence 43 from Patent W00188124.
AX421707
AX421707.1 GI:21525089
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15; Conserv
                                                                                  Method and reagent for the inhibition of erg Patent: WO 0188124-A 43 22-NOV-2001; RIBOZYME HARRMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED Location/Qualifiers
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Location/Qualifiers
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Sequence 924 from Patent
AX263533
AX263533.1 GI:16512332
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                                                                                                                                          Jarvis, T., von Carlowitz, I., Mcswiggen, J.A., Mclaughlin, F.G. Randi, A.M.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 93.8%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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/mol_type="unassigned RNA"
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AX498863
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                                                                                                                                                       Aeomica,
                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                          /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                           Location/Qualifiers
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/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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RESULT 58 AX498865

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AX735884
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AX731832/c
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Matches 15; Conserv
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Sequence 172
AX498865
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                       Homo sapiens (human)
                                                                          Sequence 1474 from Patent AX735884
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Sequences involved in phenomena of tumour suppression, tum
reversion, apoptosis and/or virus resistance and their use
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Sequence 3466 from Patent
AX731832
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Mammalia; Eutheria; Primates;
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Location/Qualifiers
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Catarrhini; Hominidae;
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  Euteleostomi; 
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RESULT 62
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BD254134.1 GI:33063904
JP 2002541795-A/1927.
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Sequences involved in phenomena of tumour suppression, reversion, apoptosis and/or resistance to viruses and thereof as medicaments
Patent: WO 03025177-A 1474 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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synthetic construct
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BD088673.1 GI:22634283
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JP 2001321190-A/917
20-NOV-2001
12-MAR-2001 JP 2001068285
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C12N15/09,C12N15/09,C12M1/00,C12Q1/68,G01N33/53,G01N33/566,
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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BD254409/c
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C12P21/02,
PC
C12P21/02,C12P21/02/A61K31/711,(C12N5/10,C12R1:91),(C12P21/02,C12R1:91),
PC
C12R1:91),
PC
C(12P21/02,C12R1:91),(C12P21/02,C12R1:91),C12N15/00,C12N5/C
PC
A61K37/02,
PC
(C12N5/00,C12R1:91)
PC
(C12N5/00,C12R1:91)
CC
Regulation of repressor genes using nucleic acid molecules
Key
Source
1.17
/organism='Eukaryote'.
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Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic acid molecules

Patent: JP 2002541795-A 1927 10-DEC-2002;

RIBOZYME PHARMACEUTICALS INC

OS Eukaryote
PN JP 2002541795-A/1927

PD 10-DEC-2002

PD 10-DEC-2002

PD 10-DEC-2002

PD 10-DEC-2002

PF 11-APR-2090 UP 2000611654

PF 11-APR-2000 UP 2000611654

PR 12-APR-1999 US 60/123990

PI LAWRENCE BLATT MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN

C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N15/10, PC
                                                                                                                                                                                                                                 OS Eukaryote
PN dp 2002541795-A/2202
PD 10-DEC-2002
PD 10-DEC-2000 dp 2000611654
PF 11-APR-2000 dp 2000611654
PR 12-APR-1999 ds 60/129390
PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC
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C12P21/02,C12P21/02/A61K31/711,(C12N5/10,C12R1:91),(C12P21/02,C12R1:91),
PC (C12P21/02,C12R1:91),(C12P21/02,C12R1:91),C12N15/00,C12N5/0
PC A61K37/02,
PC (C12N5/00,C12R1:91)
PC (C12N5/00,C12R1:91)
PC (C12N5/00,C12R1:91)
PC Regulation of repressor genes using nucleic acid molecules
Key
Location/Qualifiers
FT source 1, 17
Pr source 1, organism='Eukaryote'.
                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 17)
Blatt, I., Zwick, M., Pavco, P. and Mcswiggen, J.
Regulation of repressor genes using nucleic ac
Patent: JP 2002541795-A 2202 10-DEC-2002;
RIBOZYME PHARMACEUTICALS INC
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JP 2002541795-A/2202.
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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; Pred. No. 57;
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RESULT 65
AX216654
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3 Blatt,L., Zwick,M., Pavco,P. and Mcswiggen,J.
Regulation of repressor genes using nucleic acid
Patent: JP 2002541795-A 7194 10-DEC-2002;
RIBOZYME PHARMACEUTICALS INC
OS ENKaryote
PN JP 2002541795-A/7194
PD 10-DEC-2002
PF 11-APR-2000 JP 2000611654
                                                                                                                                                                                   1783 CGGAGGAGGCGGCA 1796
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synthetic construct synthetic construct artificial sequences
                                                                 Sequence 2096 from Patent AX216654
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BD259401
BD259401.1 GI:33069171
                                                    AX216654.1
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12-APR-1999 US 160/129390
12-APR-1999 US 160/129390
1AWRENCE BLATT,MICHAEL ZWICK,PAMELA PAVCO,JAMES MCSWIGGEN
2N15/09,A61K38/00,A61K48/00,A61P43/00,A61P43/00,C12N5/10, PC
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(C12N5/00,C12R1:91)
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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AX216927
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Matches 14
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Method and reagent for the modulation and diagnosis
nogo gene expression
Patent: WO 0159103-A 2096 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrenc
McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
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Blatt,L., Mcswiggen,J. and Chowrira,B.M. Method and reagent for the modulation and dinnogo gene expression Patent: WO 0159103-A 2369 16-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, McSwiggen, James (US); Chowrira, Bharat M.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Blatt,L., Mcswiggen,J. and Chowrira,B.M. Method and reagent for the modulation and dinnogo gene expression Patent: WO 0159103-A 2368 16-AUG-2001; RIBOZIME PHARMACEUTICALS, INC. (US); Blatt, McSwiggen, James (US); Chowrira, Bharat M. Location/Qualifiers
                                                                                                              synthetic construct
synthetic construct
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                                                                                               artificial sequences.
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/db_xref="taxon:32630"
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/db_xref="taxon:32630"
/note="Nucleic Acid"
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Sequence 569 from Patent W00188124.
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Sequence 568 from Patent WO0188124.
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                                                       Method and reagent for the inhibition of Patent: WO 0188124-A 569 22-NOV-2001; RIBOZYME PHARMACEUTICALS, INC. (US) ; GLA LOCALION/Qualifiers
                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Method and reagent for the inhibition of erg Patent: WO 0188124-A 568 22-NOV-2001; RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
                                                                                                                           Jarvis, T., von Carlowitz, I., Mcswiggen, J.A., Mclaughlin, F.G. and
                                                                                                                                                                                                      Homo sapiens (human)
                                                                                                                                                                                                                                  AX422233.1 GI:21525615
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euceleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                 Randi,A.M
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Conservative 0;
/organism="Homo sapiens"
/mol_type="unassigned RN/
/db_xref="taxon:9606"
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/mol_type="unassigned RNJ
/db_xref="taxon:9606"
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/mol_type="unassigned
/db_xref="taxon:32630"
/note="Nucleic Acid"
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Pred. No. 57;
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Query Match Best Local Similarity

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Score 14; Pred. No.

DB 1; 57;

Length 17;

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AX759707/c
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A56661
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Best Local Similarity 100.0%;
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                                                                                                                                  Patent: EP 0739898-A 28 30-OCT-1996; HOECHST AG (DE)
Other publication CZ 9600743 961016
Other publication CN 1138588 961225
Other publication PL 313207 960916
Other publication MP 8259579 961008
Other publication NO 961006 960916
Other publication NO 961006 960916
Other publication AU 4802896 960926
Other publication DE 19508923 960919.
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                                                                                                                                                                                                                                                                                                                                                        unidentified unidentified unclassified.
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Sequence 28 from Patent EP0739898.
A56661
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telerman, A., Amson, R. and Tuijnder, M.
Telerman, A., Amson, R. and Tuijnder, M.
Sequences involved in tumoral suppression, tumoral reversion,
Sequences involved in tensistance phenomena and their use as
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                  Similarity
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
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Location/Qualifiers
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                                                                                                                        ocation/Qualifiers
              0.5%;
88.2%;
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A80382
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AR045385/c
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Sequence 178
AR045385
AR045385.1
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Sequence 28 from Patent EP0726274.
A80382
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Stinchcomb,D.T., Draper,K., McSwiggen,J. and Jarvis,T.
C-myb ribozymes having 2'.5'-linked adenylate residues
Patent: US 5817796 178 06-OCT-1998;
Location/Qualifiers
1. .17
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Peyman, A.D. and Uhlmann, E.D.
G-CAP STABILIZED OLIGONUCLEOTIDES
Patent: EP 0726274-A 28 14-AUG-1996;
HOECHST AG (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         unidentified
                    AR045387 17 bp D
Sequence 180 from patent US 5817796.
AR045387
AR045387.1 GI:5966852
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178 from patent US 5
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/mol_type="unassigned_DNA"
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1. .17
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1. .17
                                                                                                                                                                                                    /organism="unknown"
/mol_type="unassigned DNA"
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AR045391/c
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1 (bases 1 to 17)

1 (bases 1, Draper,K., McSwiggen,J. and Jarvis,T. Stinchcomb,D.T., Draper,K., McSwiggen,J. and Jarvis,T. C-myb ribozymes having 2'-5'-linked adenylate residues Patent: US 5817796-A 180 06-OCT-1998;

Location/Qualifiers
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Stinchcomb,D.T., Draper,K., McSwiggen,J. and Jarvis,T.
Stinchcomb,D.T., Daving 2'-5'-linked adenylate residues
C-myb ribozymes having 2'-5'-linked adenylate residues
Patent: US 5817796-A 182 06-OCT-1998;
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Unclassified.
                                                                                                                     1 (bases 1 to 17)
Stinchcomb,D.T., Draper,K., McSwiggen,J. and Jarvis,T.
C-myb ribozymes having 2'-5'-linked adenylate residues
Patent: US 5817796-A 184 06-OCT-1998;
Location/Qualifiers
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Sequence 184 from patent
AR045391
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Sequence 28
AR111785
AR111785.1
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Peyman, A., Uhlmann, B., Breipohl, G. and
Phosphonomonoester nucleic acids proces
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Stinchcomb, D.T., Draper, K., McSwiggen, J.
C-myb targeted ribozymes
Patent: US 5646042-A 178 08-JUL-1997;
Location/Qualifiers
1 (bases 1 to 17)
Stinchcomb,D.T., Draper,K.,
C-myb targeted ribozymes
                                         Unknown.
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I52437
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Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Sequence 184 from patent US:
152443
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Stinchcomb, D.T., Draper, K., McSwiggen
C-myb targeted ribozymes
Patent: US 5646042-A 182 08-JUL-1997;
Location/Qualifiers
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Stinchcomb, D.T., D
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C-myb targeted ribozymes
Patent: US 5646042-A 184 08-JUL-1997;
Location/Qualifiers
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nilarity 88.2%;
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Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 62;
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AR188352
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AR189953
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AR189953
AR189953.1
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Ecker,J., Rothenberg,M., Lehman,A. and Roman,G.
Plant genes for sensitivity to ethylene and pathogens
Patent: US 565053-A 30 22-UUL-1997;
Location/Qualifiers
1. .17
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Sequence 30 from patent US
157029
157029.1 GI:2477442
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or concretated to levels of vascular endothelial growth factor Patent: US 6346398-A 3840 12-FEB-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unknown.
Unknown.
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                                  Unclassified.

(bases 1 to 17)

Pavco,P., McSwiggen,J., Stinchcomb,D. and Method and reagent for the treatment of direlated to levels of vascular endothelial Patent: US 6346398-A 5441 12-PEB-2002;
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/mol_type="unassigned DNA"
            Location/Qualifiers
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/mol_type="unassigned DNA"
/organism="unknown"
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AR286296
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(bases 1 to 17)

Beigelman, L., Burgin, A., Beaudry, A., Karpeisky, A., Matulio-Adamic, J., Sweedler, D. and Zinnen, S. Synthetic ribonucleic acids with RNAse activity Patent: US 6528640-A 668 04-MAR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -ب
                                                                                                                                                                                                            i (bases 1 to 17)
Pavco,P., McSwiggen,J.A., Stinchcomb,D.T. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 1607 20-WAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unknown.
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Sequence 1607 from patent
AR324205
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Sequence 667
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/mol_type="unassigned RNA"
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/mol_type="unassigned RNA"
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17 bp RNA from patent US 6617438.
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AX215457
AX215457.1 G
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Patent: WO 0159103-A 899 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, McSwiggen, James (US); Chowrira, Bharat M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 17)
Beigelman,L., Burgin,A.B., Beaudry,A., Karpeisky,A.,
Matulic-Adamic,J., Sweedler,D. and Zinnen,S.
Oligoribonucleotides with enzymatic activity
Patent: US 6617438-A 667 09-SEP-2003;
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Blatt, L., Mcswiggen, J. and Chowrira, B.M. Method and reagent for the modulation and nogo gene expression Patent: WO 0159103-A 2358 16-AUG-2001;
                                                                           synthetic construct synthetic construct
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/mol_type="unassigned RNA"
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/mol_type="unassigned RNA"
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/noTe="Nucleic Acid"
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Patent: WO 0159103-A 2361 16-AUG-2001;
RIBOZYME HARMAGEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
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                                                     Blatt,L., Mcswiggen,J. and Chowrira,B.M.
Method and reagent for the modulation and die
nogo gene expression
Patent: WO 0159103-A 2362 16 AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                                                                      Sequence 2362 from Patent AX216920
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Method and reagent for the modulation and diagnosis of cd20 and
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/mal_type="unassigned RNA"
/mal_type="trans=2630"
/mbe="Nucleic Acid"
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/mb xref="texom:32630"
/noTe="Nucleic Acid"
/organism="synthetic construct"
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AX216921
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1759 AAGATGAAGATGAGGAG 1775
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Patent: WO 0159103-A 2363 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); BlaMcSwiggen, James (US); Chowrira, Bharat Location/Qualifiers
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                                                                                                                           nogo gene expression
Patent: WO 0159103-A 2462 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
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/db_xref="taxon:32630"
/note="Nucleic Acid"
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|mol type="unassigned RNA"
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|noTe="Nucleic Acid"
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AX263537.1 GI:16512336
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 211 from Patent
AX474990
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Sequence 212 from Patent
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                    Homo sapiens (human)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Zhang,J.

Zhang,J.

Human kidney tumor overexpressed membrane protein Patent: WO 0224750-A 212 28-MAR-2002;
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/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Human kidney tumor overexpressed membrane protein 1
Patent: WO 0224750-A 629 28-MAR-2002;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and their use as
                                                                                                                                                                                                                                                                                              Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins
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Location/Qualifiers
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/mol_type="unassigned DNA"
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AX728971
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Best Local Similarity 88.2%;
Matches 15; Conservative
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Sequence 605 from Patent WO03025175.
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Molecular
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Sukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens (human)
Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                        AX759206
Sequence 2527 from Patent
AX759206
AX759206.1 GI:32253822
                                                                                                                                                                                                                                                                                                                                                                                                                           Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shannon, M., Gu, Y. and Nguyen, C.T.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 62
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Pred. No. 6:
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Oligonucleotide.
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Sequences involved in tumoral suppression, tumoral reversion, apoptosis and/or viral resistance phenomena and their use as
                                                     Epplen, J.T.

Process for the detection of restriction polymorphisms in eukaryotic genomes patent: Ep 0266787-A 11 11-MAY-1988;

Max-Planck-Gesellschaft zur Foerderung de
                                                                                                                         artificial sequences.
1 (bases 1 to 15)
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Location/Qualifiers
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
organism="synthetic construct"
/mol_type="unassigned DNA"
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88.2%; Pred. No. 6;
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1 (bases 1 to 15)
Epplen,J.T.
Process for the detection of restriction fr
polymorphisms in eukaryotic genomes
polymorphisms in 2 11-MAY-1988;
Patent: EP 0266787-A 12 11-MAY-1988;
Max-Planck-desellschaft zur Foerderung der
Location/Qualifiers
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A12052
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Harada, K., Martin, S.S. and Frankel, A.
In vivo selection of RNA-binding peptides
Patent: US 5834184-A 16 10-NOV-1998;
Location/Qualifiers
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AR053182
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larity 93.3%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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tive 0; Mismatches
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                                                                                                                                                                                                                                                      1765
           Unclassified.

1 (bases 1 to 15)
1 (bases 1 to 15)
Stinchcomb, D.T., Draper, K.G. and McSwiggen, J.
Rel a targeted ribozymes
Patent: US 5658780-A 33 19-AUG-1997;
Location/Qualifiers
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                                                                                                                                                 Sequence
161479
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Watanabe, K.A., Ren, W.-Y. and Weil, R.
Complementary DNA and toxins
Patent: US 5652350-A 19 29-JUL-1997;
Location/Qualifiers
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Sequence 19 from patent US
158728
158728.1 GI:2477966
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Watanabe, K.A., Ren, W.-Y. and Weil, R.
Complementary DNA and toxins
Patent: US 5571937-A 19 05-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unknown
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Similarity 93.3%;
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33 from patent US 5658780.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                    93.3%;
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AX635911/c
           FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protein encoded by polynucleic acid respiratory syndrome virus (PRRSV).
BD137831
                                                                                                                                                                                                                                                                                                 Paul, P.S. and Zhang, Y.
Protein encoded by polynucleic acid of porcine reproductive respiratory syndrome virus (PRRSV)
Patent: JP 2002504317-A 116 12-FEB-2002;
IOWA STATE UNIVERSITY RESEARCH FOUNDATION INC, AMERICAN CYANA OS Artificial Sequence
ON Artificial Sequence
PN JP 2002504317-A/116
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JP 2002504317-A/116.
synthetic construct
synthetic construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3050 from Patent AX635911
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14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        artificial sequences.
1 (bases 1 to 15)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genes
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                                                                                                                                                    12-FEB-2002
08-FEB-1999 JP 2000530103
06-FEB-1998 US 09/019793
PREM S PAUL, YANJIN ZHANG
C12N15,09,A61K39/12,A61P31/14,C07K14/08,C12Q1/68//C07K16/10,
                                                                                   Description of Artificial Location/Qualifiers
                                                                                                                                            C12N15/00
/organism='Artificial Sequence'
Location/Qualifiers
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/db_xref="taxon:32644"
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                                                                                                                                                                                                                      Vascular adhesion molecule and BD268813.1 GI:33078581 JP 2002537837-A/8.
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Imhof, B.A. and Lions, M.A.
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JP 2002537837-A/6
12-NOV-2002
13-MAR-2000 UP 2000603370
11-MAR-1999 EP 99200746.8
BEAT ALBEAT IMHOF MICHEL AURRAND LIONS
C12N15/09,A61K38/00,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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93.3%; Pred. No. 52;
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Vascular adhesion molecules and modulation of their function Patent: WO 0053749-A 6 14-SEP-2000;
IMHOF BEAT ALBERT (CH) ; AURRAND LIONS MICHEL (CH) ; RMF DICTIONS OF THE CONTRACT (CH) ; RMF DICTIONS OF THE CONTRA
                                                                                                                                  synthetic construct synthetic construct
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Tessier-Lavigne,M., Serafini,T., Kennedy,T.,
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11-MAR-1999 EP 99200746.8

BEAT ALBEAT IMHOF, MICHEL AURRAND LIONS

C12N15/09, AG1K38/00, AG1K39/395, AG1K39/395, AG1K39/
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/mol_type="unassigned DNA"
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/mol_type="genomic DNA"
/db_xref="taxon;32630"
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Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae;
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Sequence 8 from Patent WO0053749.
AX036056
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                                                                                        transcriptase gene
Patent: WO 02055741-A 406 18-JUL-2002;
INNOGENETICS N.V. (BE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vascular adhesion molecules and modulation of their function Patent: WO 0053749-A 8 14-SEP-2000; IMHOF BEAT ALBERT (CH); AURRAND LIONS MICHEL (CH); RMF DICES (CH)
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                                                                                                                             de Smet,K. and Stuyver,L.
Method for detection of drug-induced
                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                    entivirus group
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                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                       /organism="Human immunodeficiency
/mol_type="unassigned DNA"
/db_xref="taxon:12721"
                                                                Location/Qualifiers
1. .15
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/mol_type="unassigned DNA"
/mol_type="unassigned DNA"
/db_xref="texon:32630"
/note="primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="primer used for detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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0; Mismatches
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patent: JP 2002543802-A 3 24-DEC-2002;
Patent: JP 2002543802-A SANTE ET DE LA RE
INSTITUT NATIONAL DE LA SANTE ET DE LA RE
INSITE VISION INC
OS Artificial Sequence
PN JP 2002543802-A/3
PD 24-DEC-2002
PP 04-MAY-2000 JP 2000616394
PR 07-MAY-2000 JP 2000616394
PR 07-MAY-199 US 60/133224
PR 07-MAY-199 US 60/133224
PR 07-MAY-199 US 60/133224
PR 07-MAY-199 US LOCATION/Qualifiers
PR C12N15/09,C12Q1/68,C12N15/00
CC 01igonuclectide
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FT misc_binding (1) (16)
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Viruses; Retroid viruses; Retroviridae; Lentivirus;
lentivirus group.
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Patent: WO 02055741-A 407 18-JUL-2002;
INNOGENETICS N.V. (BE)
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synthetic construct
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Method for detection of drug-induced mutations
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1 (bases 1 to 16)
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                         /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:12721"
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Sequence 11
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AR234405
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Phillips,M.I. and Zhang,Y.
Antisense compositions targeted to .beta.1-adrenoceptor-specific mRNA and methods of use
Patent: US 6489307-A 110 03-DEC-2002;
Location/Qualifiers
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Barber, J.R., Welch, P.J., Tritz, R., Y.
Heparitis C Virus ribozymes
Patent: US 6458567-A 59 01-OCT-2002;
                                                                                                                                                                                                                                                                                             Unknown.
Unclassified
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synthetic construct
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/mol_type="genomic
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Patent: WO 0068429-A 3 16-NOV-2000;
INSTITUT MATIONAL DE LA SANTE ET DE LA
(FR); INSITE VISION INCORPORATED (US)
Location/Qualifiers
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Sequence 31 from Patent EP1076099.
AXI39183
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Mycobacterium tuberculosis
Bacteria, Actinobacteria; Actinobacteria; Actinomycetales;
Bacteria, Actinobacteria; Actinobacterium; Mycobacterium
Corynebacterineae; Mycobacteriaceae; Mycobacterium; Mycobacterium
tuberculosis complex.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic construct
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Beger, C., Barber, J. and Wong-Staal, F.
Brca-1 regulators and methods of use
Patent: WO 0170982-A 178 27-SEP-2001;
Immusol Incorporated (US); Beger, Carmela (DE)
                                                                                                                                                                                                                                                                                 Sequence 178 from Patent
AX255757
                                                                                                                                                                     synthetic construct synthetic construct
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/mcl_type="unassigned DNA"
/db_xref="taxon.32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mycobacterium tuberculosis"
|mol type="unassigned DNA"
|db xref="taxon:1773"
|noce="capture"
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AX259673/c
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AX259676/c
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Sequence 36 from Patent W00173118.
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AX259673.1 GI:16508769
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Sequence
AX259676
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Patent: WO 0173118-A 36 04-OCT-2001;
LGC (Teddington) Limited (GB)
Location/Qualifiers
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                                                                                                                        French,D.J., Mcdowell,D.G. and Brown,T. Hybridisation beacon and method of rapid sequence
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39 from Patent WO0173118.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Description of Combined DNA/RNA
Molecule:PROBE-PROBE"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/organism="synthetic construct"
/mol type="unassigned DNA"
/db xref="taxon:32630"
/note="Description of Combined DNA/RNA
Molecule:PROBE-PROBE"
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Best Local Similarity
Matches 14; Conserv
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Mycobacterium tuberculosis
Bacteria; Actinobacteria; Actinomycetales;
Corynebacteriuneae; Mycobacteriaceae; Mycobacterium; Mycobacterium
tuberculosis complex.

1 (bases 1 to 16)
Suruki, S., Nishida, M. and Takenishi, S.
Suruki, S., Nishida, M. and Takenishi, S.
Diagnosis kit of tubercle bacillus
Patent: JP 2001103981-A 31 17-APR-2001;
NT OS Mycobacterium tuberculosis
NAL NISSHINBO IND INC.SYSTEM RESEARCH CO LTD
ON Mycobacterium tuberculosis
PN JP 2001103981-A/31
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Antisense compositions targeted to _g(b)
mrna and methods of use
Patent: WO 0204623-A 110 17-JAN-2002;
University of Florida (US)
Location/Qualifiers
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Sequence 110 from Patent
AX382376
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BD013467
                                                                                                                  PD 17-APR-2001

PF 26-JUL-2000 JP 2000225985

PI SADAHIKO SUZUKI, MICHIO NISHIDA, SOICHIRO TAKENISHI PC

C12N15/09, C12N15/09, C12N1/00, C12Q1/68//(C12Q1/68, C12R1:32), I

(C12Q1/68, C12R1:325), (C12Q1/68, C12R1:33), C12N15/00, C12N15/00
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                                                                           source
                                            /organism='Mycobacterium tuberculosis'. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="unidentified"
/mol type="unassigned DNA"
/mb xref="taxon:32644"
/noTe="SYNTHETIC OLIGONUCLEOTIDE"
/organism="Mycobacterium tuberculosis"
/mol_type="genomic DNA"
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RESULT 134
A89294/c
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A88025
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Brysch, W. and Schlingensiepen, K.

ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD
PAtent: WO 9833904-A 173 06-AUG-1998;
BIOGNOSTIK GES (DE), BRYSCH WOLFGANG (DE)
LOCATION/Qualifiers
14 bp D
Sequence 1442 from Patent WO9833904.
                                                                                                                                                                                                                                                                             unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown
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Sequence 173 from patent US (
AR307962
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llarity 87.5%;
Conservative (
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                                                                                                                                                  /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
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Pred. No. 1.3e+02;
0; Mismatches 2;
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Pred. No. 64;
0; Mismatches
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ACCESSION
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A89992/c
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Similarity 92.9%;
13; Conservation
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Brysch, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD
PATENT: WO 9833904-A 1442 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
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A89294.1
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Brysch, W. and Schlingensiepen, K.

AN ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD

Patent: WO 9833904-A 1614 06-AUG-1998;

BIOGNOSTIK GES (DE), BRYSCH WOLFGANG (DE)
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          1 (bases 1 to 14)
Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonucleotide preparation method Patent: EP 0856579-A 173 05-AUG-1998;
BIOGNOSTIK GES (DE)
                                                                                                                                        Sequence 173
A89992
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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 Location/Qualifiers
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Pred. No. 64;
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Pred. No. 64;
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AX239937
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ACCESSION
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Sequence 27 from Patent WO0236761.
AX587251
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Modified Oligonucleotides for mismatch discrimination
Patent: WO 0164958-A 64 07-SEP-2001;
Epoch Biosciences, Inc. (US)
Location/Qualifiers
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                                                                                                                    Patent: WO 0236761-A 27 10-MAY-2002;
DANA FARBER CANCER INSTITUTE (US)
Location/Qualifiers
                                                                                                                                                                       D'Andrea,A.D., Taniguchi,T., Timmers,C. and Grompe,M. Methods and compositions for the diagnosis of cancer susceptibilities and defective dna repair mechanisms and
                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
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/mol_type="unassigned DN7
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                       /mol_type="unassigned
/db_xref="taxon:9606"
1. 14
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                                      /note="Intron/Exon Junction of FANCD"
                                                                                          organism="Homo sapiens"
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Primates;
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Pred. No. 64;
O; Mismatches
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BD066807/c
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         AUTHORS Schlingenstepen,K.H. and Brysch,W.

FITLE An antisense oligonuclectide preparation method patent: JP 2001511000-A 1442 07-AUG-2001;

BIOGNOSTIK GESSELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH DI JP 2001511000-A/1442

PD Unknown JP 2001511000-A/1442

PD 07-AUG-2001

PF 30-JAN-1998 JP 1998532533

PF 31-JAN-1998 JP 1998532533

PF 31-JAN-1997 EP 9710157
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Schlingensiepen K.H. and Brysch, W.

An antisense oligonucleotide preparation method

Patent: JP 2001511000-A 173 07-AUG-2001;

E BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH

OS Unknown

PN JP 2001511000-A/173

PD 07-AUG-2001

PF 30-JAN-1998 JP 1998532533

PF 30-JAN-1998 JP 1998532533

PR 31-JAN-1998 JP 1998532533

PR 31-JAN-1997 EP 97101531.8

PI KARL HERMANN SCHLINGENGIEPEN, WOLFGANG BRYSCH

PC C12N15/11, C077H21/04, A61K31/70

CC An antisense oligonucleotide preparation method FH Ke
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An antisense oligonucleotide
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Unknown

JP 2001511000-A/1442
07-AUG-2001
30-JAN-1998 JP 1998532533
31-JAN-1997 EP 97101531.8
KARL HEMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
C12N15/11,C07H21/04,A61K31/70
An antisense oligonucleotide preparation method
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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BD137809
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AUTHORS
AND AUTHOR
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synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 14)

1 (bases 1 to 14)

Paul, P.S. and Zhang, V.
Protein encoded by polymucleic acid of porcine
respiratory syndrome virus (PRRSV)
Patent: JP 2002504317-A 94 12-FEB-2002;
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An antisense oligonucleotide
BD066979
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BD137809
BD137809.1 GI:23232754
JP 2002504317-A/94.
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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                                                  l Similarity
13; Conserv
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Patent: JP 2002504317-A 98 12-FEB-2002;
IOWA STATE UNIVERSITY RESEARCH FOUNDATION INC, AMERICAN CYANAMID OS Artificial Sequence
PN JP 2002504317-A/98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BD137813 14 bp
Protein encoded by polynucleic a
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1 (bases 1 to 14)
Paul, P.S. and Zhang, Y.
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                      CCACCCCTTAACC 2040
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Artificial Sequence
JP 2002504317-A/94
                                                                                                                                                                                                                                  06-FEB-1998 US 09/019793

PREM S PAUL, YANJIN ZHANG
C12N15/09, A61K39/12, A61P31/14, C07K14/08, C12Q1/68//C07K16/10,
                                                                                                                                                                                                                                                         12-FEB-2002
08-FEB-1999 UP 2000530103
06-FEB-1998 UP 09/0197
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06-FEB-1998 US 09/019793
PREM S PAUL, YANJIN ZHANG
C12N15/09,A61K39/12,A61P31/14,C07K14/08,C12Q1/68//C07K16/10,
                                                  Conservative
                                                                                                                                                                                  source
                                                                                                                                                                                                          Description of Artificial Sequence: Synthetic DNA
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                  /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                             92.9%;
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                                                 Score 12.4; D
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RESULT 144

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synthetic construct
artificial sequences.

1 (bases 1 to 14)
Paul, P.S. and Zhang, Y.
Protein encoded by polynucleic acid of porcine reproductive and respiratory syndrome virus (PRRSV)
Patent: JP 2002504317-A 111 12-FEB-2002;
IOWA STATE UNIVERSITY RESEARCH FOUNDATION INC, AMERICAN CYANAMID CO
                                                                                                                                                                                      Paul, P.S. and Zhang, Y.

Protein encoded by polynucleic acid of porcine respiratory syndrome virus (PRRSV)

Patent: JP 2002504317-A 113 12-FEB-2002;
IOWA STATE UNIVERSITY RESEARCH FOUNDATION INC, I
                                                                                                                                                                                                                                                                                                                                             14 bp DNA linear PAT 18-: Protein encoded by polynucleic acid of porcine reproductive respiratory syndrome virus (PRRSV).
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BD137826
BD137826.1 GI:23232771
JD 2002504317-A/111.
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1 (bases 1 to 14)
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JP 2002504317-A/113.
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06-FEB-1998 US 09/019793

PREM S PAUL, YANJIN ZHANG

C12N15/09, A61X39/12, A61P31/14, C07K14/08, C12Q1/68//C07K16/10,

C12N15/09
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llarity 92.9%;
Conservative
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JP 2002504317-A/111
12-FEB-2002
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06-FEB-1998 US 09/019793
PREM S PAUTIN ZHANG
C12N15/09,A61K39/12,A61P31/14,C07K14/08,C12Q1/68//C07K16/10,
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JP 2002504317-A/113
12-FEB-2002
                                                           Description of Artificial
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Location/Qualifiers
/organism='Artificial
Location/Qualifiers
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                                                                                                                       Method and reagent for molecule participating BD201812
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein encoded by polynucleic acid of porcine respiratory syndrome virus (PRRSV) Patent: JP 2002504317-A 120 12-FEB-2002; IOMA STATE UNIVERSITY RESEARCH FOUNDATION INC, FOR Artificial Sequence PN JP 2002504317-A/120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       respiratory syndrome BD137835
Sukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 14)
PavCo,P.A., Roberts,E., Jarvis,T., Coeshott,C. and Mcswiggen,J.A. Method and reagent for treating diseases or conditions concerning
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                                                                               Homo sapiens (human)
                                                                                                              BD201812.1 GI:33011582
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1 (bases 1 to 14)
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                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Description of Artificial Sequence:Synthetic DNA Location/Qualifiers
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06-FEB-1998 US 09/019793
PREM S PAUL, YANJUN ZHANG
C12N15/09, A61K39/12, A61F31/14, C07K14/08, C12Q1/68//C07K16/10,
                                                                                                                                                                                                                                                                                                                                                                                                  /organism='Artificial
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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92.9%;
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Pred. No. 64
                                                                                                                                 14 bp RNA linear PAT 17-JUL-200 treating diseases or conditions concerning in vasculogenic response.
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I (bases 1 to 14)

II (bases 1 to 14)

Blatt, L. Mcswiggen, J.A., Roberts, E., Pavco, P.A. and Macejak, D.

TLE Enzymatic nucleic acid treatment of diseases or conditions related to hepatitis C virus infection

RNAL Patent: JP 2002512791-A 3038 08-MAY-2002;

RIBOZYME PHARMACEUTICALS INC

OS Hepatitis virus (hepatitis C virus)

PD 08-MAY-2007

PD 08-MAY-2007

PF 0-MAY-2007
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                                                                                                                     PAVCO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid treatment to hepatitis C virus infection.
                                                                                                                                                                                                                      PF 26-APR-1999 JP 2000545991

27-APR-1998 US 60/100842 PR

27-APR-1999 US 09/257608,23-MAR-1999 US 09/274553 PI

1AWRENCE BLATT, JAMES A MCSWIGGEN, ELISABETH ROBERTS, PAMELA A PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C12N15/09,A61K31/7088,A61K31/7125,A61K48/00,A61P3/10,A61P17/06, PC
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27-MAR-1998 US 60/079678
PAMELA A PAVCO,ELISABETH ROBERTS,THALE JARVIS,CLAIRE COESHOTT,
JAMES A MCSWIGGEN
                                                                                                                                     C12N9/00,A61K31/7105,A61K38/21,A61K48/00,A61P31/12,C12N15/09,A61K37/66,
C12N15/00
                                                                 Enzymatic nucleic acid treatment related to hepatitis C virus infection. hepatitis C virus infection/Qualifie Key
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/organism='Hepatitis virus virus)'
Location/Qualifiers
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/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Pred. No. 64;
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                                                                   Location/Qualifiers
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A90173/c
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Best Local Similarity 92.9%;
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1 (bases 1 to 15)

Brysch, W. and Schlingensiepen, K.

AN ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD
PACENT: WO 9833904-A 354 06-AUG-1998;
BICGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
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13; Conserv
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Sequence 354 from Patent WO9833904.
A88206
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Sequence 354 from Patent EP0856579.
A90173
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Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonucleotide preparation method Patent: EP 0856579-A 354 05-AUG-1998;
                                                                                                                                                                                                                      unidentified
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 CATCTGTGAGCTGC 1
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                                                                                     /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="genomic RNA"
/db_xref="taxon:32644"
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Pred. No. 78;
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Pred. No. 64;
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Pred. No. 78;
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AR113878/c
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Ribozyme treatment of diseases or conditions related to levels intercellular adhesion molecule-1 (ICAM-1)

Patent: US 6132967-A 324 17-OCT-2000;

Location/Qualifiers
                                                                                        15 bp I
Sequence 247 from patent US 5567809.
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Sequence 324 from patent US
AR056120
AR056120.1 GI:5981697
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Sequence 324 from patent US 6132967.
AR113878 GI:14094200
1 (bases 1 to 15)
Apple,R.J., Erlich,H.A.,
Methods and reagents for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Intercellular adhesion molecule-1 (ICAM-1) ribozymes
Patent: US 5837542-A 324 17-NOV-1998;
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 15)
Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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 Griffith,R.L. and Scharf,S.J. HLA DRbeta DNA typing
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Sequence
152067
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Swiggen,J.A. and Mamone,J.Anthony.
Hammerhead ribozymes for preferred targets
Patent: US 5646020-A 9 08-JUL-1997;
Location/Qualifiers
                                                                                                                                                      1 (bases 1 to 15)
Stinchcomb,D.T., Draper,K.G. and McSwiggen,J.
Rel a targeted ribozymes
Patent: US 5658780-A 201 19-AUG-1997;
Location/Qualifiers
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/mol_type="unassign
                                                                                                                        organism="unknown"
/mol_type="unassigned DNA"
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AR179965
AR179965.1
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Sequence 23 from Patent WO0114531.
AX088088
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Vogelstein,B., Kinzler,K.W., Zhang,L.
Gene expression profiles in normal and
Patent: US 6333152-A 391 25-DEC-2001;
Location/Qualifiers
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Vogelstein,B., Kinzler,K.W., Zhang,L. and Zhou,W.
Gene expression profiles in normal and cancer cells
Patent: US 6333152-A 33 25-DEC-2001;
Location/Qualifiers
                                   May,G.D., Kmiec,E.B. and Rice,M.C.
Cell-free assay for plant gene targeting and conversion
Patent: WO 0114531-A 23 01-MAR-2001;
The Samuel Roberts Noble Foundation, Inc. (US)
Location/Qualifiers
                                                                                                                          synthetic construct synthetic construct
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organism="synthetic construct"/mol_type="unassigned DNA"
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The Samuel Roberts Noble Foundation, Inc. Location/Qualifiers
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13; Conserv
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Musa sp.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopaida; Zingiberales; Musaceae;
                                                                                                                                          May,G.D., Kmiec,B.B. and Rice,M.C. Cell-free assay for plant gene targeting a Patent: WO 0114531-A 25 01-MAR-2001; The Samuel Roberts Noble Foundation, Inc.
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Spermatophyta; Magnoliophyta; Liliopsida;
clade; Panicoideae; Andropogoneae; Zea.
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/note="Plasmid pTsm153
                                                                                /organism="Musa sp."
/mol_type="unassigned DNA"
/db_xref="taxon:46838"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Zea mays"
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AX088091
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Nicotiana tabacum
Nicotiana tabacum
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Trachec
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
asterids; lamiids; Solanales; Solanaceae; Nicotiana.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core ev
asterids; lamiids; Solanales; Solanaceae; Nicotiana.
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The Samuel Roberts Noble foundation, Inc. (US)
Location/Qualifiers
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27 from Patent WO0114531.
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AX636058/c
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SOURCE
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unidentified unclassified.
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Eukaryota; M
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stinchcomb, D.T., Dudycz, L.W., Chowrira, B., Grimm, S., Direnzo, A., Karpeisky, A., Draper, K.G., Kisich, K., Matulic-Adamic, J., Mcswiggen, J.A., Modak, A., Pavco, P., Beigelman, L., Sullivan, S.M., Sweedler, D., Thompson, J.D., Tracz, D., Usman, N., Wincott, F.E. and
                                                                                     AX636058
Sequence 3197 from Patent
AX636058
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    unidentified unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 359 from Patent AX633220
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                          unidentified
                                                                                                                                                                                                                                                                                                                                                                                      Fatent: EF 1260586-A 359 27-NOV-2002; RIBOZYME PHARMACEUTICALS, INC. (US)
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                                                                        AX636058.1
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Similarity 92.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                             /mol_type="unassigned RNA"
/db_xref="taxon:32644"
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                                                                        GI:28471672
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92.9%; Pred. No. 78;
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prognose or treat epilepsy
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BD061455
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Sequence 468 from Patent EP1288296
AX711168
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                                        synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 15)
1 (bases 1 to 15)
1 Ishibashi, K. and Tsuji, A.
Method for selectively separating living
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Herpes simplex virus unknown type
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
specific gene
Patent: JP 2001286285-A 17 16-OCT-2001;
LABORATORY OF MOLECULAR BIOPHOTONICS
                                                                                                                                        BD061455.1 GI:22607061 JP 2001286285-A/17.
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Similarity 92.9%;
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/mol_type="unassigned RNJ
/db_xref="taxon:32644"
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                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Herpes simplex virus
/mol_type="unassigned RNA"
/db_xref="taxon:126283"
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An antisense oligonucleotide preparation method
L Patent: JP 2001511000-A 354 07-AUG-2001
BICGNOSTIK GESELLSCHAFT FUR BICMOLEKULARE DIAGNOSTIK MBH
OS Unknown
PN JP 2001511000-A/354
PD 07-AUG-2001
PF 30-JAN-1998 JP 1998532533
PF 30-JAN-1998 JP 1998532533
PF 31-JAN-1997 EP 97101531.8
PI KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
PC C12N15/11, C07H21/04, A6LK31/70
CC An antisense oligonucleotide preparation method FH KA
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An antisense oligonucleotide pl
BD065719
BD065719.1 GI:22611322
JP 2001511000-A/354.
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PN JP 2001
PD 16-CCT-
PF 28-APR-
PI KANAME
PC C12N15/
G01N33/566,
PC G01N33/566,
PC G01N33/566
PC C12N5/
CC Probe
FH Key
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15 bg MBD104667 m 15 bg Kit and method for determining BD104667 BD104667.1 GI:22650241
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                                                                                                                                                                                                                                                                                                                                         30-JAN-1998 JP 1998532533
31-JAN-1997 EP 97101531.8
XARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
C12N15/11, C07H21/04, A61K31/70
An antisense oligonuclectide preparation method
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KANAME ISHIBASHI,AKHIKO TSUJI
C12N15/09,C12N1/02,C12N5/10,C12Q1/68,G01N33/48,G01N33/53,
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JP 2001286285-A/17
16-OCT-2001
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                                                                                                                                                                                                                                                                                   /organism='Unknown'.
Location/Qualifiers
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                                                                                                                                                                                                                                                       synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 15)
                                                                                                                            NISHIDA
OS Art
PN WO
PD 06-
PF 01-
PR 01-
PR 111
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PN WO 01
PN 06-DB
PD 06-DB
PP 01-JU
PF 11D-TU
PI HIDET
MATSUMURA,
PI SHOGO
PC C12Q1
CC C12Q1
CC C12Q1
CC CSC
FH Key
FT SOURC
                  MATSUMURA,
PI SHOGO
PC C12Q1
CC Descr
FH Key
FT sourc
                                                                                                                                                                                                                                 Nishida,M.

Kit and method for determining HLA type

Rit and method for determining HLA type

Patent: WO 0192572-A 889 06-DEC-2001;

NISSHINBO INDUSTRIES INC, SYSTEM RESEARCH INC, HIDETOSHI IN

KAGIYA, TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO MORIYA,
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13; Conserv
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Rit and method for determining BD104785

BD104785.1 GI:22650359
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Patent: WO 0192572-A 771 06-DEC-2001;
NISSHINBO INDUSTRIES INC, SYSTEM RESEARCH INC, HIDETOSHI INOKO,
KAGIYA, TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO MORIYA, MICH
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synthetic construct
artificial sequences.
1 (Dases 1 to 15)
Inoko,H., Kagiya,T., Ichihara,T.,
Nishida,M.
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ilarity 92.9%;
Conservative
                                           SHOGO MORIYA, MICHIO NISHIDA
C12Q1/68, C12M1/00, C12N15/09, G01N33/53
Description of Artificial Sequence: capture
Key Location/Qualifiers
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w0 0192572-A/771
06-DEC-2001
01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
HIDETOSHI INOKO,TAEKO KAGIYA, TATSUO ICHIHARA, YOSHIYUKI
                                                                                                                       Artificial Sequence
WO 0192572-A/889
WO-DEC-2001
01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
HIDETOSHI INOKO,TAEKO KAGIYA,TATSUO
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C12Q1/68,C12M1/00,C12N15/09,G01N33/53
Description of Artificial Sequence:capture
Key Location/Qualifiers
/organism='Artificial
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 78;
0; Mismatches
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YA, MICHIO
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BD167992/c
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AR307963/c
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Matches 13; Conserv
                                                                                                                              ORGANISM
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                                                                                                                                                                                                                                                                                        16
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MITSUBISHI CHEMICAL CORP, TORU TSUJI, HIROSHI YANAGAWA
OS Artificial Sequence
PN WO 0226964-A/39
PD 04-APR-2002
PF 26-SEP-2001 WO 2001JP008387
PF 26-SEP-2001 WO 2001JP008387
PR 27-SEP-2000 JP 00P 293692,06-FEB-2001 JP 01P 029138
TORU TSUJI, HIROSHI YANAGAWA
PC C12M15/09,C12P21/02
CC Description of Artificial Sequence:Synthesized FH Key
                                                                                                                                                                                      Sequence 174 from patent AR307963
                                                                                                                                                                                                                                                                                                                                             l Similarity
13; Conserv
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                                        Antisense modulation of raidd expression Patent: US 6551826-A 174 22-APR-2003; Location/Qualifiers
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synthetic construct
artificial sequences.
1 (bases 1 to 16)
                                                                                      1 (bases 1 to 20) Watt, A.T.
                                                                                                                              Unknown
                                                                                                                                               Unknown.
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WO 0226964-A/39.
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                                                                                                                                                                                                                                                                                        GCTGCAGGGTCAGGC 2
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larity 86.7%; Pred. No. 1.2e+02;
Conservative 0; Mismatches 2
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
/organism="unknown"
/mol_type="genomic DNA"
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AX023424/c
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AR010038/c
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AR034773/c
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Best Local Similarity 69.6%;
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Sequence 51
AR034773
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                     AX023424
Sequence 39 from Patent
AX023424
AX023424.1 GI:10183824
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Johnson, E.M. and Bergemann, A.D.
Monoclonal antibodies to the pur protein
Patent: US 5869622-A 51 09-FEB-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown.
Unclassified.
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Location/Qualifiers
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AR010038
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Johnson, E.M. and Bergemann, A.D.
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Similarity 69.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="unknown"
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Page 42
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SOURCE	synthetic construct
	artificial sequences.
REFERENCE	1
AUTHORS	Lipford, G.B., Heeg, K. and Wagner, H.
TITLE	G-motif oligonucleotides and uses thereof
JOURNAL	Patent: WO 0014217-A 39 16-MAR-2000;
	LIPFORD GRAYSON B (DE) ; HEEG KLAUS (DE) ; WAGNER HERMANN (DE) ;
	CPG IMMUNOPHARMACEUTICALS GMBH (DE)
FEATURES	Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Variation
                                                                                                                                                                                                                                                                                                                                                                                     (WHED ) WHITEHEAD INST BIOMEDICAL RES
                       1823
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                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGCCGGCGAGGTGGAGCTCTC
 GGCCGGCGAGGTGGAGCTCTC
                                                                                          21
                     0.8%;
conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibitor
                                                                                          B₽;
                                                                                                                                                                                                                                                                                                                                                                Ireland
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                                                                                          N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers replace(11,A) /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /standard_name= "single nucleotide polymorphism"
                                                                                          Α,
                                                                                                                                                                                                                                                                                                                                                                 JS,
                                                                                          ر:
و ر:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymorphism containing DNA fragment #155.
                                                                                                                                                                                                                                                                                                                                                                 Lander ES
                                                                                            10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                        Score 21; pred. No.
                                             0,
                                                                                            <u>ი</u>
                       1843
 21
                                                                                            ω
                                              Mismatches
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H
                                                                                            0
                                                                                           U; 0 Other;
                                                         58
                                                                    1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           science;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer;
                                               ..
                                                                    Length 21;
                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; inflammation; ds.
                                               0
                                               Gaps
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RESULT 4
AAT32789/c
ID AAT32789 standard; DNA; 25
XX
AC AAT32789;
XX
DT 18-FEB-1997 (first entry)
XX
Triple helix-forming oligo
XX
Triple helix; triplex form
KW purification; double-stran
XX
Synthetic.
XX
PN W09618744-A2.
XX
PD 20-JUN-1996.
XX
XX
OS-NOV-1995; 95WO-FR0014
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Double-stranded (ds) DNA can be purified from complex mixtures of nucleic acids, proteins, endotoxins, nucleases, etc. by passing the mixture over a support to which an oligonucleotide is covalently attached; the oligonucleotide is able to form a triple helix by hybridisation with a specific sequence present in the dsDNA. The present sequence is a preferred oligonucleotide which can form a triple-helix with the homopurine target sequence in AAT32776. The target sequence may be present naturally, e.g. in a plasmid origin of replication, or can be introduced artificially. The method is particularly suited to
                                                                                                                                                                      Triple helix-forming oligonucleotide for purifying plasmid pXL2725
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        purification of plasmid DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Page 26; 34pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9618744-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Triple helix; triplex formation; Hoogsteen base pairing; plasmid; purification; double-stranded DNA; homopyrimidine; polypurine; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hybridising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RHON ) RHONE POULENC RORER SA.
                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                          1771
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            helix-forming
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                                                                                                                                                                                                                                                                                                                                                          AGGAGGAGGAGGCGAGGC 1792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of double stranded DNA by triple helix formation - comprises ng immobilised oligo-nucleotide to specific sequence in target
                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 2 A; 14 C; 1 G; 8 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Scherman
                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                             triplex formation; Hoogsteen base pairing; plasmid, double-stranded DNA; homopyrimidine; polygurine; p)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94FR-00015162.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95WO-FR001468
    95WO-FR001468
                                                                                                                                                                                                    entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                     95.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                      Score 20.4; I
Pred. No. 11;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        complex mixtures of nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                               polypurine; pXL2725;
                                                                                                                                                                                                                                                                                                                                                                                                                      25;
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RESULT 5
AAS19343/c
ID AAS19343;

XX AAS19343;

XX AAS19343;

XX Oligonuclectide sequ
XX Synthetic.

XX Synthetic.

XX Expent_region 5...
FT repeat_region 5...
FT repeat_unit 5...
FT repeat_unit 5...
FT repeat_unit 9/rpt
FT CCOUZET J, Schermar
XX 26-MAY-2000; 2000US-
PA (AVET) AVENTIS PHAF
XX 27 AVET 3 AVENTIS PHAF
XX WPI; 2002-097772/13.
XX Purifying double-str
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Best Local S
Matches 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 25 BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-300660/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide sequence used to purify plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Page 18; 34pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Purificn. of double stranded DNA by triple helix formation - c
hybridising immobilised oligo-nucleotide to specific sequence
   Purifying double-stranded (ds)
                                    WPI; 2002-097772/13.
                                                                                                                              26-MAY-2000; 2000US-00580923
                                                                                                                                                             25-MAY-2001; 2001WO-US017122
                                                                                                                                                                                                                                                                                                                                                                                                           ss; DNA purification; triple helix; plasmid purification; XL2725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence into plasmid pBKS+)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Double-stranded (ds) DNA can be purified from complex mixtures of nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RHON ) RHONE
                                                                                                 (AVET ) AVENTIS PHARMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1771 AGGAGGAGGAGGCGAGGAGGC 1792
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scherman
                                                                  Scherman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         POULENC RORER SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94FR-00015162.
                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers 5. .25
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                                                                                                                                                                                                                                                                                           /rpt_type=
                                                                                                                                                                                                                                                       /note= "CCT repeat type"
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                                                                                                 SA.
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                                                                  Wils
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 20.4;
Pred. No. 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ש
                                                                  Ψ,
      DNA from a solution containing dsDNA and
                                                                  Blanche
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
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                                                                     Cameron
                                                                                                                                                                                                                                                                                                                                                                                                                                             XL2725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 6
AAL615
XX
AAL615

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ikappab r; antisense; immune response; inf tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          other components, comprises passing the solution comprising a covalently coupled oligonucleotide shelix with the dsDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130469.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 7.2; Page 20; 40pp;
  13-NOV-2001; 2001US-00993731
                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                        05~NOV-2002; 2002WO-U8035597
                                                                                                                                                                      WO2003042360-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1771 AGGAGGAGGAGGCGAGGC 1792
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S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; r; antisense; immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry
                                                                                                                                                                                                                                                                                          /*tag= \u03b4
/mod_base= OTHER
/note= "2'-methoxyethyl (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprises
                                                                                                                                                                                        /*tag= c
/mod_base= OTHER
/note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
1. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "Phosphorothioate backbone; All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Score 20.4; D; ; Pred. No. 11; 0; Mismatches
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                                                                                                                                                                                                                              (2'-MOE)
                                                                                                                                                                                                                                                                                                                                         (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     n through a support
able to form a triple
                                                                                                                                                                                                                              nucleotides
                                                                                                                                                                                                                                                                                                                                      nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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AAL61563/c
ID AAL615
XX
AC AAL615
AC AAL615
AC Human
XX
Human
XX
Human
XX
Humour
XX
Humour
XX
Homo s
OS Synthe
CS Synthe
FT modifi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to antisense compounds targetted to a nucleic acid conclecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, ITKBR, I-kappa-B-related, ikappabr, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                          Human, inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy, tumour; prophylaxis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighteimmune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
 WO2003042360-A2.
                                                                        modified_base
                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                         Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                       22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL61563;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL61563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Page 74; 108pp;
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                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                               sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; S. larity 100.0%; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                               /*tag= a
/mod_base= OTHER
/note= "Phosphorothioate b
/note= "Phosphorothioate b
are 5-methylcytidines"
           /*tag= c
/mod_base= OTHER
/note= "2'-methoxyethyl
                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                            note=
                                                                                                                        *tag=
                                                                             . 20
                                                                               _base= OTHER
e= "2'-methoxyethyl
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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7.3:
                                  (2'-MOE) nucleotides
                                                                                            (2'-MOE)
                                                                                                                                                                       backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Other;
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                                                                                            nucleotides"
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                                                                                                                                                                       cytidine residues
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RESULT 8

AAL61578/c

ID AAL61578 standard; DNA; 20 BE

XX

AC AAL61578;

XX

AC AAL61578;

XX

CAL61578;

XX

AC AAL61578;

XX

DE Human inhibitor-kappa B-R ant

XX

Human; inhibitor-kappa B-R; I

KW Human; inhibitor-kappa B-R; I

KW ikappab r; antisense; immune

KW tumour; prophylaxis; phosphoz

XX

Homo sapiens.

OS Synthetic.

XX

Key

FT modified_base 1, 20

FT modified_base 2, mod_base= 0, mod_base=
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130503.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1323 GGGGACCTCTTCTCCAAGGC 1342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Page 74; 108pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                       prophylaxis; phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGGGACCTCTTCTCCAAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Watt AT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 4 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001US-00993731
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                                                                        /*tag= a
/mod_base= OTHER
/mote= "Phosphorothioate backbone;
/note= "Phosphorothioate backbone;
are 5-methylcytidines"
                                                                                                                                                                                                                                                                          Location/Qualifiers
      base= OTHER
== "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                             58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
7.3;
            (2'-MOE) nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
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                                                                                                                                                            All
                                                                                                                                                         cytidine
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ARESULT 9
AAL64/c
ID AAL615
XX AAL615
AC AAL615
XX 22-SEP
XX 22-SEP
XX Human
XX Human;
KW Human;
KW ikappa
KW tumour
XX
XX Homo s
OS Synthe
XX Key
FH Key
FH Key
FH Key
FH Key
                                                                                                                                                                                                                                                                                                                                                             片
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to antisense compounds targetted to a nucleic acid cc molecule encoding human inhibitor-kappa B-R (also known as I-KappaBR, CC IKBR, I-kappa-B-related, ikappabr, nuclear factor of kappa light college gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful cfor treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving concerned cytokine expression, or a result of infection (e.g. bacterial, college or parasitic). They are useful for diagnostics, therapeutics, componing or parasitic) in the expression of cormation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide crargetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 20
                                                                                                                        Human; inhibitor-kappa B-R; I-kappaBR; ikappab r; antisense; immune response; tumour; prophylaxis; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighteimmune response or infection.
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                                                                                                                                                                                                                     22-SEP-2003
                                                                                                                                                                                                                                                     AAL61546
                                                                                                                                                                                                                                                                                 AAL61546 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
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                               modified_base
                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                       1621 CGCTCAGCTGTGCTCAGCAG 1640
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                                                                                                                                                                                      inhibitor-kappa
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                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 100.0%;
20; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 75; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                               CGCTCAGCTGTGCTCAGCAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 4 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                     (first entry)
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/*tag= c
/mod_base= (
/*tag= a
/mod_base= OTHER
                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note= "2'-methoxyethyl (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                         B-R
                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 20; DB
100.0%; Pred. No. 7.
ive 0; Mismatches
                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                      antisense oligonucleotide,
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                                                                                                                                           IKBR; I-kappa-B-related; NFKBIL2;
infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20
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                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 10
AAL61549/c
ID AAL615
XX AAL615
XX 22-SEP
XX 22-SEP
XX Human
XX Human;
KW Human;
KW ikappa
KW tumouz
XX OS Homo £
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                                                                                                                                                                                                                                                                                             S
                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                             Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappaBB-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NPKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides targeted to nucleic acids encoding inhibitor kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
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                                                                                                                           22-SEP-2003
                                                                                                                                                         AAL61549;
                                                                                                                                                                                       AAL61549 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 9 C; 4 G; 2 T; 0 U; 0
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                                                                                            inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     base
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                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "2'-methoxyethyl
16. .20
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/mod_base= OTHER
/note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Phosphorothioate backbone; All cytidine residues
                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                             Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                           7.3;
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                                                                                                                                                                                                                                                                                                                                                                                           Other;
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                                                                                                                                                                                                                                                                                                                                                            Length 20;
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RESULT 11
AAL61554/c
ID AAL61554 standard; DNA; 20 B
XX
AC AAL61554;
XX
DT 22-SEP-2003 (first entry)
XX
DE Human inhibitor-kappa B-R an
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                                                                                                                                                                                                                  Best Local
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
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modified_base
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20; Conserv
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                                                                                                                                                              GTTCATGGAGAGCGAGTGCT
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                                                                                                                                                                                                                  0.8%; Score 20; DB llarity 100.0%; Pred. No. 7. Conservative 0; Mismatches
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/mod_base=
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/note= "2'-methoxyethyl
16. .20
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are 5-methylcytidines"
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= "2'-methoxyethyl (2'-MOE)
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Human inhibitor-kappa B-R antisense oligonucleotide,

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Best Local S
Matches 20
                                                                                                                                      The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
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ikappab r; antisense; immune response;
                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 74; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC
                            1213
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20
                                                                     Similarity
               CCATCTGTCAGAACCTCCAG 1232
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                                                        Conservative
                                                                                                                BP; 4
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
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/mod base OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
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/mod_base= OTHER
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                                                                     0.8%;
                                                                                                                3 C; 8 G; 5 T; 0 U; 0 Other;
                                                        0; Mismatches
                                                                     Score 20;
Pred. No.
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infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (2'-MOE) nucleotides
                                                        0
                                                                                   Length 20
                                                        Indels
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RESULT 12 AAL61561/c ID AAL615

AAL61561 standard; DNA; 20

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                            Matches
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Best Local
                                                                                                      The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-ike 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research respents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
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                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
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1300 CCATGGTCATCTGTGAGCAG 1319
                           l Similarity
20; Conser
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                                                                                 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                           Conservative
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/mod_ba
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are 5-methylcytidines"
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Pred. No.
                            Mismatches
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infection; inflammation; therapy;
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                                        DB 1;
7.3;
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                                                   The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
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b r; antisense; immune response; infection; inflammation;
; prophylaxis; phosphorothicate; ss.
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/note= "Phosphorothicate
are 5-methylcytidines"
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-NOE)
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/note= "2'-methoxyethyl
16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            All cytidine residues
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Sequence 20

BP; 4

A; 5 C;

J

G; 4 T; 0 U; 0 Other;

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RESULT 14

ARL61540/c
ID ARL615
XX Human;
XX Ikappa
XX Exprise
XX Synthe
XX Synthe
YFT modifi
FT modifi
FT
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Best Local S
Matches 20
The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I kappaBR, IXBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAL61540 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
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                                                                                                                                                                                                                                                                                                                                                                                       New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-NOV-2002; 2002WO-US035597.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003042360-A2
                                                                                                                                                                                                                                                                                                                                     Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-NOV-2001; 2001US-00993731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003-468635/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inhibitor-kappa B-R antisense oligonucleotide, ISIS #130465.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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                                                                                                                                                                                                                                                                                                                                     Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCTCTTCTCCAAGGCAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTCTTCTCCAAGGCAGGAG 1347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                     74; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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/mod_base=
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/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note= "2'-methoxyethyl (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .3;
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RESULT 15
AL61545/c
ID AAL615
XX AAL615
XX AAL615
XX AAL615
XX AL6165
XX AL6165
XX AL6165
XX AL6165
XX AL6161
XX Human;
XX Human;
XX Ikappa
XX Exprise modifi
FT Tmodifi
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                                                                                                                                                                                        New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAL61545
  The invention relates to antisense compounds targetted to a nucleic aci molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-NOV-2001; 2001US-00993731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-2002; 2002WO-US035597
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                                                                                                                                               Claim
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                                                                                                                                                                                                                                                                                                                2003-468635/44.
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                                                                                                                                            Page 74; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTGTGCAACGATTACTTCAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of various members of a biological pathway. They are also antisense therapy. The present sequence is an oligonucleotide to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                 Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%;
ilarity 100.0%;
Conservative (
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/mod_base= OTHER
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= "2'-metho:
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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                                                                                                                                                                                                                    heightened
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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AAL61562
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Synthetic.
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                                            New antisense oligonucleotides targeted to nucleic acids encodininhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heiginmune response or infection.
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o r; antisense; immune response;
; prophylaxis; phosphorothioate;
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/note= "2'-methoxyethyl
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are 5-methylcytidines"
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7.3;
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:ion; inflammation;
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Best Local Similarity Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                       modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human;
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                                                   Monia BP,
                                                                                                                                             05-NOV-2002; 2002WO-US035597.
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                    WPI; 2003-468635/44
                                                                                                             13-NOV-2001; 2001US-00993731
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b r; antisense; immune response; infection; inflammation; therapy;
                                                                                ISIS
                                                 Watt AT,
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                                                                                  PHARM INC
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE)
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/note= "Phosphorothicate backbone; All cytidine residues
are 5-methylcytidines"
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1. 20
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e= "2'-methoxyethyl (2'-MOE)
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Pred. No.
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7.3;
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Sequence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;

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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa BR, polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       108pp; English.
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RESULT 18
AAL61570/c
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XX AAL615
XX AAL615
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CONTRIBUTION
CONTRIBUTI
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Best Local S
Matches 20
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20; Conserv
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/mod_ba
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/mod_base= OTHER
/note= "Phosphorothioate backbone;
/note= "Phosphorothioate backbone;
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/mod_base= OTHER
/mote= "2'-methoxyethyl
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Pred. No.
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                                                                                                                                                                                                                                                                        (2'-MOE)
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13-NOV-2001; 2001US-00993731

05-NOV-2002; 2002WO-US035597

22-MAY-2003

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RESULT 19
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    molecule encoding human inhibitor-kappa B-R (also known as I-kappabR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBL12) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-468635/44.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130496.
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                                  WO2003042360-A2
                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ikappab
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b r; antisense; immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prophylaxis; phosphorothicate; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%;
(larity 100.0%;
Conservative
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1. .20
                                                                                                                                                                                                                     /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine
are 5-methylcytidines"
                                                                                                                                                                                                         note=
                                                                                         note=
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                                                                                                                                                                                . 20
                                                                                                                        base= OTHER
                                                                                         "2'-methoxyethyl
                                                                                                                                                                                                            "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Score 20; DB
%; Pred. No. 7.3
0; Mismatches
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7.3;
                                                                                         (2'-MOE)
                                                                                                                                                                                                            (2'-MOE)
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                                                                                            nucleotides"
                                                                                                                                                                                                            nucleotides"
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RESULT 20
AAL61555/c
ID AAL615
XX AAL615
XX AAL615
XX Luman
XX Human
XX Human
XX Human
XX Human
XX Homo s
OS Synthe
XX Synthe
FI Ry
FI modifi
FI modifi
FI modifi
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IXBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides targeted to nucleic acids encoding inhibitor kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
                                                                                                                                                                                                                                    Human, inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; ikappab r; antisense; immune response; infection; inflammation; tumour; prophylaxis; phosphorothicate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 74; 108pp;
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                                                         modified_base
                                                                                                                                       modified_base
                                                                                                                                                                                        Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                    Human inhibitor-kappa B-R antisense oligonucleotide,
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 modified_base
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20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTGCGCCACTATGAGGAGGA 1504
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                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                              /mod_base= OTHER
/note= "Phosphorothioate
are 5-methylcytidines"
1 . . 5
 /note=
16. .20
                                                                                                                                                         Location/Qualifiers
                                mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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                "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7.3;
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                (2'-MOE)
                                                                                              backbone; All cytidine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                  nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                       ISIS #130480.
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RESULT 21
AAL61550/c
ID AAL615
XX AAL615
XX AAL615
XX AAL615
XX Human
XX Human
XX Human
XX Human
XX Homo s
OS Synthe
XX Synthe
XX Synthe
XX Modifi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                          Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
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                                                                                                                                                                                                                          Human inhibitor-kappa B-R antisense oligonucleotide,
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                                                               modified_base
                                                                                                                Synthetic
                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                         (first entry)
/*tag= a
/mod_base= OTHER
/mote= "Phosphorothicate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                Location/Qualifiers
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 20; 7.3;
                     backbone; All cytidine residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 22
AAL61551/c
ID AAL61551 standard; DNA; 20
XX AAL61551;
XX
DT 22-SEP-2003 (first entry)
XX
DE Human inhibitor-kappa B-R
XX
M ikappab r; antisense; immm
XM ikappab r; antisense; immm
XM tumour; prophylaxis; phosp
XX
OS Homo sapiens.
OS Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to antisense compounds targetted to a nucleic acid compounds encoule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IXBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for disgnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                              Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 7 A; 7 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; Page 74; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotides targeted to nucleic acids encoding inhibitor kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-NOV-2001; 2001US-00993731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-2002; 2002WO-US035597
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                                                                                                                  Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-468635/44.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                          1099 AGTGCTGCGTGGTTATTGCA 1118
                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                               20;
                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                AGTGCTGCGTGGTTATTGCA 1
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                                                                                                                                                 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note=
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                                                                                                                                                                                                                                                                                                                                                        100.0%;
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= "2'-methoxyethyl (2'-MOE)
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                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                              Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 20 7.3;
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                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                      ISIS #130476.
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RESULT 23
AAL61566/c
ID AAL615
XX
AC AAL615
XX
AC AAL615
XX
DT 22-SEP
XX
DE Human
XX
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                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                  The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa BR, polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monia BP,
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modified_base
               Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130491.
                                                                                                      AAL61566 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC
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                                             22-SEP-2003
                                                                         AAL61566
                                                                                                                                                                                                       1112
                                                                                                                                                                            20
                                                                                                                                                                                                                                   l Similarity 100.0%;
20; Conservative
                                                                                                                                                                                            TATTGCACAGGTCCTCCAAG 1131
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                                                                                                                                                                            TATTGCACAGGTCCTCCAAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Watt AT;
                                                                                                                                                                                                                                                                                           BP; 5 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
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                                             (first entry)
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are 5-methylcytidines"
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/note= "2'-methoxyethyl
/ic. .20
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                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      base= OTHER
                                                                                                                                                                                                                                                  0.8%; Score 20;
100.0%; Pred. No.
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                                                                                                      20
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                                                                                                      ВP
                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                Length 20;
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                                                                                                                                                                                                                                      Indels
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RESULT 24
AAL61577/c
ID AAL61577;
XX

standard; DNA; 20

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Query Match
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Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monia BP,
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20
                                                                                       Similarity
20; Conserv
                                                                                                                                                                                     20
GACTITCCCAGGGCAGCTGA 1
                                           GACTITCCCAGGGCAGCTGA 1366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Watt AT;
                                                                                                                                                                                     BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
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/mod_ba
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e= "2'-methoxyethyl
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                                                                                                               100.0%;
                                                                                                                                         0.8%;
                                                                                                          ; Score 20;
%; Pred. No.
                                                                                          0;
                                                                                             Mismatches
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infection; inflammation;
                                                                                                            DB 1;
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                                                                                             0
                                                                                                                                         Length 20;
                                                                                                  Indels
                                                                                               0,
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                                                                                               Gaps
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                                                   Query Match
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Matches 20
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Synthetic.
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                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                    Example 15; Page 75; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighteimmune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-NOV-2001; 2001US-00993731.
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b r; antisense; immune response; infection; inflammation; therapy;
; prophylaxis; phosphorothioate; ss.
20
                                                                    Similarity
                                                                                                            20
                           CGATGCCTACGAGCTGCTGG 1602
                                                    .larity 100.0%; F
Conservative 0;
                                                                                                           BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
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/mod_base= OTHER
/mote= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "2'-methoxyethyl
16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "Phosphorothicate
are 5-methylcytidines"
                                                                                 0.88;
                                                                    Score; Pred.
                                                      Mismatches
                                                                                 20;
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                                                                  7.3;
                                                                                 DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     backbone;
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                                                      Indels
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ARLESULT 25
ARAL6126
ID ARAL61
AC AR
                                                         The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKappaBR, IKappaBB, IKappaBR, IKappaBR, IKappaBR, IKappaBR, IKappaBR, IKappaBB, IKappaBR, IKa
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AAL61581
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
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b r; antisense; immune response; infection;
; prophylaxis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                response or infection.
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/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
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infection; inflammation; therapy;
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also
                                                                                                                                                                                      New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
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The invention relates to antisense compounds targetted to a nucleic aci molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of

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RESULT 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide
                                 New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for disgnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ikappab r; antisense; immune response;
tumour; prophylaxis; phosphorothioate;
                                                                                             WPI; 2003-468635/44.
                                                                                                                                                                  13-NOV-2001; 2001US-00993731
                                                                                                                                                                                         05-NOV-2002; 2002WO-US035597
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130507
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                                                                                                                      Monia
                                                                                                                                                                                                                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibitor-kappa B-R; I-kappaBR;
b r: antisense; immune response;
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                                                                                                                                           PHARM INC
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                             /*tag= c
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE)
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/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
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Pred. No.
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infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.3;
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Claim 3; Page 75; 108pp; English

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RRESULT 29
RAL61539/c
ID AAL615
XX AAL615
AC AAL615
XX Human
XX Human;
XX Homo s
SS Homo s
SS Homo s
SS Withe
XX Homolfi
FT modifi
FT mo
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                                                        WPI; 2003-468635/44
                                                                                                                                                                                                                13-NOV-2001; 2001US-00993731
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Synthetic:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human inhibitor-kappa B-R antisense oligonucleotide, ISIS
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  New antisense oligonucleotides targeted to nucleic acids
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                                                                                                                                                                PHARM INC
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/mod_base= OTHER
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Phosphorothioate
are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "2'-methoxyethyl
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Pred. No.
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7.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (2'-MOE)
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     encoding
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RESULT 30
AAL61547/c
ID AAL615
XX AAL615
XX AAL615
XX AAL615
XX Human;
XW Human;
XW Human;
XW Ikappa
KW Ikappa
KW Ikappa
KW Iwoux
XX
FH Key
FT modifi
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Matches 20
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                                  13-NOV-2001;
                                                                       05-NOV-2002;
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   (ISIS-) ISIS
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b r; antisense; immune response; infection; inflammation; therapy;
; prophylaxis; phosphorothioate; ss.
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20; Conserv
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   PHARM INC
                                      2001US-00993731.
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/mod_base= OTHER
/note= "2'-methoxyethyl
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/note= "Phosphorothicate backbone;
are 5-methylcytidines"
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                                                                                                                                                                                base= OTHER
== "2'-methoxyethyl (2'-MOE)
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RESULT 31
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Synthetic.
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Similarity 100.0%; Pred. No.
20; Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                      /*tag= b
/mod_base= OTHER
/note= "2'-methoxyethyl (
16. 20
                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "Phosphorothioate backbone;
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                   *tag=
                                                                          base= OTHEK
5= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                  OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 20, 7.3;
                                                                                                                                                                                     (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
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                                                                                                                                                                                                                                                                                                     All cytidine residues
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22-MAY-2003

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RESULT 32
AL61575/c
ID AL615
XX AL615
XX AL615
XX Human
XX Human;
KW Ikappa
KW Ikappa
KW Ikappa
KW Ikappa
KW Ikappa
KW Ikappa
KW Kumour
XX
KW Ikappa
KW Kumour
XX
KW Ikappa
KW I
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Best Local
                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides targeted to nucleic acids encoding inhibitor kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL61575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL61575 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-NOV-2001;
                                    modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibitor-kappa B-R antisense oligonucleotide, ISIS #130500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; br; antisense; immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       r; antisense;
prophylaxis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Page 75;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGTGCTGGAGGAGGCCAAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Watt AT;
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                                                                                                                                                                                                      /mod_base= OTHER
/note= "Phosphorothioate backbone;
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                              /mod_base= OTHER
/note= "2'-methoxyethyl
                                                                                                                                           *tag=
                                                                                                                                                                                                                                                                                                                    *tag=
                                        . 20
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           a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
7.3;
                                                                     (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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                                                                                                                                                                                                                                              All cytidine
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AAL61584/c
ID AAL615
XX
AC AAL615
XX 22-SEP
XX Human
XX Human;
XW ikappa
XW ikappa
XW tumour
XX Homo s
OS Synthe
XX Synthe
XX Homo s
XX 
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa Bpa Ight polypeptides gene enhancer in B-cells inhibitor-like 2 and NPKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                           Key
modified_base
                                                                                                                                                                                                                                                                                                                      Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r_i antisense; immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                   Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides targeted to nucleic acids encoding inhibitor kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAL61584;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; Page 75; 108pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1530 CTGGAGGAGGCCAAGACCTG 1549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
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                                                                                                                                                                                                                                                                                              prophylaxis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTGGAGGAGGCCAAGACCTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
   /mod_base= OTHER
/note= "Phosphorothioate
are 5-methylcytidines"
                                                                                   /*tag= a
                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'mod_base= OTHER
'note= "2'-methoxyethyl (2'-MOE) nucleotides"
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                                                                                                                                                                                                                                                                                        phosphorothicate;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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7.3;
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                              backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20
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                              All
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                              cytidine residues
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RESULT 34
AAL61585/c
ID AAL615
XX AAL615
XX AAL615
XX AAL615
XX AAL615
XX Human
XX Human;
XX S Synthe
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                Homo sapiens.
Synthetic.
                                                            ikappab r; antisense; immune response;
tumour; prophylaxis; phosphorothioate;
                                                                              Human; inhibitor-kappa B-R; I-kappaBR;
ikappab r; antisense; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-NOV-2002; 2002WO-US035597.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                         Human inhibitor-kappa
                                                                                                                                                           22-SEP-2003
                                                                                                                                                                                          AAL61585
                                                                                                                                                                                                                       AAL61585 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-NOV-2001; 2001US-00993731.
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                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                TGCAGCTGAGGCTGCAGCCC 1709
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                       (first entry)
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/*tag=
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16. .20
/*tag=
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'note= "2'-methoxyethyl
                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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_e= "2'-
                                                                                                                           B-R antisense oligonucleotide, ISIS #130510.
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                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                           Score 20; DB 1; Pred. No. 7.3;
                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                            IKBR; I-kappa-B-related; NFKBIL2;
infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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AAL61552/c
ID AAL61552/c
XX AAL6155
AC AAL6155
XX AAL6155
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XX LUMAN 1
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XX HUMAN;
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Best Local :
                                                                                                                                                                                                                                                                                                                                                         molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, I-Kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NREBL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for disgnosties, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and indistinguishing between functions of various members on a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2;
                                Human inhibitor-kappa B-R antisense oligonucleotide,
                                                              22-SEP-2003
                                                                                              AAL61552
                                                                                                                           AAL61552 standard; DNA;
                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 6 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to antisense compounds targetted to a nucleic a molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-468635/44.
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                                                                                                                                                                                                                                      1730 AACCAGACTGCGGGAGCTCA 1749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Page 75; 108pp; English.
                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Watt AT;
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                                                              (first entry)
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/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                  0.8%; Score 20;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                    DB 1; Length 20; 7.3;
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                                 ISIS #130477.
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RESULT 36
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AC AAL615
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Best Local
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Synthetic.
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tumour; prophylaxis; |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
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     AAL61553;
                                                  AAL61553 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-468635/44.
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                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                         20 BP; 4 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002WO-US035597
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1. .20
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/note= "Phosphorothicate backbone; All cytidine residues
are 5-methylcytidines"
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/mod_base= OTHER
/note= "2'-methoxyethyl
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/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
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phosphorothicate; ss.
                                                                                                                                                                                                                                                                                       0.8%; Score 20;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                          7.3;
                                                                                                                                                                                                                                                                                                              DB 1;
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                                                                                                                                                                                                                                                                   Gaps
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, ITKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                        Query Match
Best Local S
Matches 20
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modified_base
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                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Page
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inhibitor-kappa B-R, useful for diagnosing or treating diseases
associated with expression of inhibitor-kappa B-R, e.g., a heigh
immune response or infection.
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1194 AAGCCTGTGCAGAGGGCAGC 1213
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                                                                   Similarity
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                                                                                                                                 BP; 2 A; 8 C; 5
                                           Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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are 5-methylcytidines"
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note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= b
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e= "2'-methoxyethyl (2'-MOE) nucleotides"
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(၂)
                                                              Score 20; ; Pred. No.
                                             Mismatches
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                                                                                                                                   U; 0 Other;
                                                                   DB 1
7.3;
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                                                                                      1.
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tion; inflammation; therapy;
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                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleotides'
                                               Indels
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RESULT 37
AAL61556/c
The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                   New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
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ikappab r; antisense; immune response; infection; inflammation; therapy;
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e= "2'-methoxyethyl
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Query Match Best Local Similarity

0.8%;

Score 20; Pred. No.

DB 1; 7.3;

Length 20

Sequence 20

BP; 5 A;

9 C; 4 G; 2 T; 0 U; 0 Other;

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RESULT 38

AL61542/c
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, I-kappa-B-related, ikappab r, nuclear factor of kappa 19th polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKSII2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
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o r; antisense; immune response;
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infection; inflammation; therapy;
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappaB-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving
                                                                                                New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
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 Page 74; 108pp;

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o r; antisense; immune response; infection; inflammation;
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are 5-methylcytidines"
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7.3;
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                                                                                                                                                                                                                                                                                         (2'-MOE)
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                                                   New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height
                                                                                                                                                                                                                                                                                                                                                                        Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           increased cytokine expression, or a result of infection (e.g. bacterial viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
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/note= "Phosphorothioate backbone;
are 5-methylcytidines"
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RESULT 41
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Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; inhibitor-kappa B-R; I-kappaBR; ikappab r; antisense; immune response; tumour; prophylaxis; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human inhibitor-kappa B-R antisense oligonucleotide,
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New antisense oligonucleotides targeted to nucleic acion inhibitor-kappa B-R, useful for diagnosing or treating
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                                                                                     WPI; 2003-468635/44.
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/mod_base= OTHER
/note= "2'-methoxyethyl
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are 5-methylcytidines"
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infection; inflammation; therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r_i antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3;
                                                                                    13-NOV-2001; 2001US-00993731
                                                                                                                                       05-NOV-2002; 2002WO-US035597
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                                                                                                                                                                                                                                                                                                                                                                                                                              "2' -methoxyethyl
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pred. No.
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7.3;
                                                                                                                                                                                                                                                                                                                  (2'-MOE) nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                              (2'-MOE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotides'
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RESULT 43
AAL61573/c
ID AAL615
XX AAL615
XX AAL615
XX AAL615
XX AAL615
XX Human;
XX Human;
XX Itagpa
XX It
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; inhibitor-kappa B-R; I-kappaBR; ikappab r; antisense; immune response; tumour; prophylaxis; phosphorothicate;
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                                  22-MAY-2003
                                                                                                                                                                                                                      modified_base
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                                                                                    WO2003042360-A2
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                                                                                                                                  /*tag= (/mod_base= (
                                                                                                                                                                                                                                                                           /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "Phosphorothicate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                /mod_ba
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                                                                                                                                                                                                                                              note=
                                                                                                                                                                                                                            . 20
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e= "2'-methoxyethyl (2'-MOE)
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                                                                                                                                        -methoxyethyl
                                                                                                                                                                     OTHER
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.
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infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
. 7.3;
                                                                                                                                           (2'-MOE)
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                                                                                                                                        nucleotides
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RASSULT 44
AAL61568/c
ID AAL615
XX AAL615
XX 22-SEP
XX Human;
XW Human;
XW Human;
XW Lumour
XX Lumour
XX Homo s
OS Homo s
OS Synthe
XX Homolfi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                         modified_base
                                                                                                                                                                                                                                                                                               Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 75; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
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                                                                                  modified_base
                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                            22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                       AAL61568;
                                                                                                                                                                                                                                                                                                                                                                                  AAL61568 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 8 C; 2 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                     kappab
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                                                                                                                                                                                                            sapiens
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                                                                                                                                                                                                                                       inhibitor-kappa B-R; I-kappaBR; IKBR;
b r; antisense; immune response; infect
; prophylaxis; phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                          (first entry)
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                           /note=
16. .20
                                                                                              /mod_base= OTHER
/note= "Phosphorothicate
are 5-methylcytidines"
                                                                                                                                                                    Location/Qualifiers
                                                                   /*tag=
               *tag=
                              . 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                          _base= OTHER
e= "2'-methoxyethyl
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                a
                                                                                                                                                                                                                                                                                                                                                                                  20
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                       IKBR; I-kappa-B-related;
infection; inflammation;
ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
7.3;
                                           (2'-MOE)
                                                                                                                backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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                                           nucleotides"
                                                                                                                A11
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                                                                                                                cytidine residues
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'mod_base= OTHEF

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RESULT 45
AAL61538/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                  Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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modified_base
                                                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                 (first entry)
/*tag= a
/mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
1..5
                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 20;
100.0%; Pred. No.
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7.3;
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AAL61541 s

XX AAL61541;

XC AAL61541;

XX 22-SEP-200

XX 22-SEP-200

XX Human inhi

XX Human; inh

XX Human; i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                         Homo sapiens
Synthetic.
                                                                                                                                          Human; inhibitor-kappa B-R; I-kappaBR; ikappab r; antisense; immune response; tumour; prophylaxis; phosphorothioate;
                                                                                                                                                                                                                                                      Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                  22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTGAATGAGATGAGGACCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGAATGAGATGAGGACCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "
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== "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             base≕ OTHER
e= "2'-methoxyethyl
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              859
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 20; 7.3;
                                                                                                                                                                           infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (2'-MOE)
                                                                                                                                                                                                      I-kappa-B-related; NFKBIL2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleotides"
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                                                                                                                                                                                 inflammation; therapy;
                                                                                                                                                                                                                                                               ISIS #130466.
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., a heightened
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Location/Qualifiers

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RESULT 47
AAL61572/c
ID AAL61
XX AAL61
XX AAL61
XX AAL61
XX AAL61
XX Human
XX Human
KW Human
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                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                               The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I kappa-B-related, ikappab r, nuclear factor of kappa Iight polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
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 Human; inhibitor-kappa
ikappab r; antisense; :
                                             Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130497.
                                                                              22-SEP-2003
                                                                                                           AAL61572
                                                                                                                                        AAL61572 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 15; Page 74; 108pp; English.
                                                                                                                                                                                                                                   940 TCCTTGCGGAGCAGAACCAC 959
                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  response or infection.
                                                                                                                                                                                                                                                                              20;
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                                                                                                                                                                                                                 TCCTTGCGGAGCAGAACCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Watt AT;
                                                                                                                                                                                                                                                                                                                                        BP;
                                                                                                                                                                                                                                                                          0.8%; Solarity 100.0%; 1
Conservative 0;
                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*ttag= b
/mod_base= OTHER
/note= "2'-methoxyethyl (
                                                                                                                                                                                                                                                                                                                                        w
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/mod_base= OTHER
/mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
/note= "enethylcytidines"
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                                                                                                                                                                                                                                                                                                                                        >
                                                                                                                                                                                                                                                                                                                                       5 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           base= OTHER
   1 B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2;
immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "2'-methoxyethyl (2'-MOE) nucleotides'
                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                           Score 20; Pred. No.
                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                           DB 1;
7.3;
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                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                         Length 20;
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                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                              Gaps
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RESULT 48
AAL61583/c
ID AAL615
XX
AC AAL615
XX

AAL61583 standard;

DNA; 20

0

AAL61583

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                                                                                                                                     The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa-light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                      Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                         New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-468635/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-NOV-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tumour; prophylaxis; phosphorothicate; ss
                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 75; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-NOV-2001; 2001US-00993731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
                           1488 CGCCACTATGAGGAGGAACT 1507
 CGCCACTATGAGGAGGAACT
                                                      0.8%; Score 20; DB ilarity 100.0%; Pred. No. 7. Conservative 0; Mismatches
                                                                                                               BP; 3 A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002WO-US035597.
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/note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note= "Phosphorothicate
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mod base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   _base= ОТНЕК
}= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER
                                                                                                             G; 6 T; 0 U; 0 Other;
                                                                    DB 1;
7.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       backbone; All cytidine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -MOE)
                                                          0
                                                                                   Length 20
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                                                                                                                                                                                                                                                                                                                                                                                                            a heightened
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                                                         Gaps
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22-SEP-2003

(first entry

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                                                                                                                                The invention relates to antisense compounds targetted to a nucleic acid complexible encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r. nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NPKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                      Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                               Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-NOV-2001; 2001US-00993731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inhibitor-kappa B-R; I-kappaBR; IXE
r; antisense; immune response; inf
prophylaxis; phosphorothioate; ss.
                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                             Page 75; 108pp; English
  CTCCATACCGTGCAGCTGAG 1
                         crccaraccerecaecreae 1699
                                                                                                           BP; 4 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002WO-US035597.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= ":
16. .20
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1. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          *tag= b
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                                                0.8%; Er
100.0%; Er
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e= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "2'-methoxyethyl
                                                                  Score 20; DB 1; Length 20; Pred. No. 7.3;
                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IKBR; I-kap
infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (2'-MOE) nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (2'-MOE)
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tion; inflammation; therapy;
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                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                        heightened
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                                                        Gaps
                                                                                                                                                                                                                                                                                                                    acid
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RESULT 49
AAL61543/c
ID AAL61543;
XX
AX
CAAL61543;
XX
CAAL
                                                            The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa 15th polypeptides gene enhancer in B-cells inhibitor-ike 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13~NOV-2001; 2001US-00993731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-NOV-2002; 2002WO-US035597
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15; Page 74; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotides targeted to nucleic acids encoding ibitor-kappa B-R, useful for diagnosing or treating diseases ociated with expression of inhibitor-kappa B-R, e.g., a heightened une response or infection.
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are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note= "2'-methoxyethyl (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note=
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infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (2'-MOE)
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tion; inflammation; therapy;
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molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBII2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inhibitor-kappa B-R, useful associated with expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense oligonucleotides targeted to nucleic acids encoding ibitor-kappa B-R, useful for diagnosing or treating diseases ociated with expression of inhibitor-kappa B-R, e.g., a heightened une response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP,
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b r; antisense; immune response; infection; inflammation; therapy;
; prophylaxis; phosphorothioate; ss.
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are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                           New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
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are 5-methylcytidines"
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.e= "2'-methoxyethyl
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The invention relates to antisense compounds targetted to a nucleic act molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa 119ht polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBII2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving inhibitor-kappa B-R such as a heightened immune response involving

increased

cytokine

expression,

a result of

infection

(e.g.

bacterial

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Example 15; Page 74; 108pp; English

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RESULT 52
AAL6155/C
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                                                                                                                                                New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130490.
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                                                                        Claim 3; Page 74; 108pp; English.
                                                                                                                                                                                                                                                  WPI; 2003-468635/44.
                                                                                                                                                                                                                                                                                                                                                                                                   13-NOV-2001; 2001US-00993731
                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-NOV-2002; 2002WO-US035597.
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                                                                                                                          response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGCAACAGCTGGAAGAGGCT 1
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                                                                                                                                                                                                                                                                                                   Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ilarity 100.0%;
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/mod_ba
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/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
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/note= "2'-methoxyethyl
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; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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The invention

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relates to antisense compounds targetted to a nucleic acid ding human inhibitor-kappa B-R (also known as I-kappaBR,

heightened

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RESULT 53
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Best Local |
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New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
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                                                                                                                                                             Monia BP,
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression: Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonuclectide targetted to human inhibitor-kappa B-R DNA
        Monia BP,
                                                                                                                                                                   05-NOV-2002; 2002WO-US035597
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130504.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inhibitor-kappa B-R; I-kappaBR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              r;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
     Watt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                 2001US-00993731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/mote= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
i..5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "2'-methoxyethyl
16..20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              *tag=
                                                                                                                                                                                                                                                                                                                             base OTHER
= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immune response;
                                                                                                                                                                                                                                                                                                                                                                                           Ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IKBR; I-kappa-B-related; NFKBIL2;
infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
7.3;
                                                                                                                                                                                                                                                                                                                                   (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                            (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                                   nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                     Claim 3; Page 75; 108pp; English.
                                                                                                                                                                                                                  WPI; 2003-468635/44.
```

for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA The invention relates to antisense compounds targetted to a nucleic act molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful acid

Sequence 20 BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;

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밁
                 S
                                          Matches
                                                    Query Match
Best Local
                    1652 CCAGCTGCAGAGGCAGGTCT 1671
20 CCAGCTGCAGAGGCAGGTCT 1
                                         Similarity
20; Conserv
                                        0.8%; Solarity 100.0%; I Conservative 0;
                                                  Score 20; DB 1;
Pred. No. 7.3;
                                         Mismatches
                                         0,
                                                             Length 20;
                                          Indels
                                         0
                                         Gaps
                                          0
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AAH62253
ID AAH
XX AAH
XX AAH
XX AAH
XX AAH
AC AAH
XX Sin
XX Sin
XX Hom
XX AAH
XX Hom
XX Hom
XX Hom
XX W2
PT Var
PT Var
PT Var
PT Var
PT I7-
PF 17-
PF 1 Car
XX WPI
XX New
PT ince
                                                                                                                                                                                                                                                                                                                                                        RESULT 55
                                                                                                                                                                                                                                                                                                                  AAH62253;
                                                                                                                                                                                                                                                                           NF-kappa-B inhibitor polymorphism containing DNA fragment #154.
                                                                                                                                                                                                                                                                                                                                     AAH62253 standard; DNA;
                                                                                                                                                                                                                                                                                               12-SEP-2001
                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                     ВP
```

Single nucleotide polymorphism; SNP; human; cancer; inflammation; heart disease; paternity testing; forensic science; ds.

Homo sapiens

```
WO200138576-A2
                                                                              Variation
                                                                            Location/Qualifiers replace(11,T)
                                 /standard_name= "single nucleotide polymorphism"
```

31-MAY-2001.

24-NOV-1999; 17-NOV-2000; 2000WO-US031639 99US-0167334P

(WHED) WHITEHEAD INST BIOMEDICAL RES

Cargill M, Ireland JS, Lander ES

WPI; 2001-367705/38

New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.

Claim 1;

Page 42;

80pp;

English.

contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in a applications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis

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RESULT 56
AAT1992A7
ID AAT199
XX AAT199
XX DT 15-AF
XX PUR 6
XX PUR 6
KW CANCE
XX SYNTH
OS HOMO
XX 10567
XX 06-JU
XX 07-JU
XX 06-JU
XX 08-JU
PR 02-FE
PR 02-FE
PR 02-FE
PR 06-JU
XX WPI;
XX Seam
XX WASS
XX Exam
XX Ex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 20
                                                            This oligonucleotide represents a dimer sequence of the upTAR was used as a competitor oligonucleotide in a gel shift assay binding of PUR protein to DNA. The PUR sequence can be used to chemical or biological compounds that bind to PUR or binding f PUR. Inhibitors of PUR activity may be used to treat hyperprol
  Sequence
                                                                                                                                                                                           Example; Col 39; 64pp; English.
                                                                                                                                                                                                                                     Assays for PUR protein ligands protein or fragments, to treat
                                                                                                                                                                                                                                                                                                      WPI; 1997-488859/45.
                                                                                                                                                                                                                                                                                                                                                  Bergemann
                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-AUG-1992;
02-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-SEP-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5672479-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PUR element;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        upTAR element
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT99287 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                            (MOUN ) MOUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
    24
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                                                                                                                                                                                                                                                                                                                                                  ₿
  BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 3 A; 9 C; 6 G; 3 T; 0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human; c-myc; inhibitor; hyperproliferative disease; ss; e; hybridisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                              as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dimer oligonucleotide for binding
                                                                                                                                                                                                                                                                                                                                                                                              SINAI SCHOOL MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                   92US-00938189.
93US-00014943.
95US-00470911.
                                                                                                                                                                                                                                                                                                                                                  Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-00486421
                                              cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry
  4
C;
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  16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,,</u>
  <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 19.4;
Pred. No. 11;
                                                                                                                                                                                                                                     or modulators - using immobilised PUR hyper-proliferative diseases, e.g. can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
  0
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  <del>..</del>
  0
  ä
  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Other;
                                                                                 PUR or binding for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ۲.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human PUR-alpha
                                                         be used to identify binding fragments of hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                             for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                          cancer.
                                                                                                                                                  and
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AAX04092
ID AAXX
XX
AC AAXX
XX
AC IDT 12-J
DT 12-J
XX
DE UPTJ
XX
KW PUR

12-APR-1999

(first

entry)

PUR element;

upTAR element oligonucleotide purine-rich probe.

PUR-alpha; hyperproliferative disease;

human;

片

Н

24

RESULT 58

AAX04092 standard;

DNA;

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RESULT 57
AAV31744
ID AAV31744
AC AAV31744
AC AAV31
AC AAC AAV31
AC AAV31
AC AAV31
AC AAC AAC AAC AAC AAC AAC AAC AAC AA
δ
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Best Local S
Matches 21
                                                                              Query Match
Best Local (
                                                    Matches
                                                                                                                                                                                               This is the nucleotide sequence of a purine-rich oligonucleotide used a competitor with the PUR element in the method of the invention, involving the use of the PUR protein and its fragments, which inhibit protein binding to PUR element or other proteins. Inhibitors of PUR activity may be useful for treating viral infections and hyperproliferative diseases such as cancer
                                                                                                                                                     Sequence 24 BP; 4 A; 4 C; 16 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                        Example 7.1.1; Col 33; 63pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 PUR protein and its fragments -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-321632/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-AUG-1992;
02-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PUR-alpha gene; inhibition; hyperproliferative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV31744;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV31744 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US5756684-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MOUN ) MOUNT SINAI SCHOOL MEDICINE
     1772 GGAGGAGGAGGCGAGGAGGCGGC 1795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1772 GGAGGAGGAGGCGGAGGAGGCGGC 1795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21;
                                                       21;
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                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                          or other proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGAGGCGGAGGCGGAGGC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ΑĐ,
                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92US-00938189.
93US-00014943.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-00470911
                                                                            0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of a purine-rich oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%;
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전
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        viral infection; cancer;
ss.
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Pred. No. 18;
0; Mismatches
                                                                              Score 19.2;
Pred. No. 18
                                                                                                                                                                                                                                                                                                                                                                          English.
                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    that inhibit PUR protein binding
                                                                                 18
                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB
                                                                                                         ۲.
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                                                                                                      Length 24;
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                                                       Indels
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                                                       0,
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                                                       Gaps
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ARESULT 59
ARZ99647
ID ARZ99
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XX 12-JU
DT 12-JU
XX 12-JU
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Matches 21
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02-FEB-1993;
06-JUN-1995;
                                                                                                                                                                                                      G-motif oligonucleotide; vaccine; Toxoplasmosis; viral infection; antigen presenting cell activation; natural killer cell; septic shock; cytotoxic T-lymphocyte; inflammation; autoimmune disease; rheumatoid arthritis; Crohn's disease; sarcoidosis; multiple sclerosis; Kawasaki syndrome; graft-versus-host disease; transplant rejection; helper T cell response 1-mediated disease; Lyme arthritis; Streptcoccal induced arthritis; chronic inflammatory bowel disease; psoriasis vulgaris; experimental allergic encephalomyelitis; insulin-dependent diabetes mellitus; bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
Homo sapie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleotide sequence of non-G-motif oligonucleotide Pur-alpha-Ori.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       binds to an epitope of the PUR protein. Antibodies that bind to the PUR protein and neutralise PUR activity may be used to treat hyperproliferative diseases such as cancer. PUR antibodies may be used diagnostically to detect aberrant expression of the PUR protein and/or mutations in the PUR gene. The present sequence represents an upTAR element oligonucleotide purine-rich probe, which is used in an example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ99647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ99647 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 24 BP; 4 A; 4 C; 16 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUN-1995;
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                              16-MAR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a monoclonal antibody that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-152881/13
                                                                               WO200014217-A2
                                                                                                                                                                                insulin-dependent dia
parasitic infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rrom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the present invention
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93US-00014943.
95US-00470911.
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Pred. No. 1
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                                                                                                                                                                                     tumour; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    specifically
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                                                                                                          conjugation. The specification describes compositions comprising G-motif coligonucleotides. The G-motif oligonucleotides inhibit activation of antigen presenting cells by inhibiting the uptake of DNA by a cell, by stimulating natural killer cells, or by co-stimulating cytotoxic T-clymphocytes. The G-motif oligonuclectides may be used for the productions of vaccines for treating septic shock, inflammation, autoimmune diseases (e.g. rheumatoid arthritis, Crohn's disease, sarcoidosis, multiple collection), helper T cell response 1-mediated disease and transplant crejection), helper T cell response 1-mediated diseases (e.g. Streptococcal induced arthritis, Lyme arthritis, chronic inflammatory complaints, and insulin-dependent disease (e.g. Compalations, parasitic infections (e.g. Leishmaniasis or Toxoplasmosis), wire lifections, parasitic infections (e.g. Leishmaniasis or Toxoplasmosis), crifections, spontaneous abortions and human immunodeficiency virus confiduce proliferation of bone marrow cells, especially macrophage confidences cells
                                      Matches
                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       septic shock, rhe virus infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Compositions comprising G-motif oligonucleotides useful for treating eseptic shock, rheumatoid arthritis, diabetes and human immunodeficiency
                                                                                                        Sequence 24 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                       The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 14; Page 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CPGI-) CPG IMMUNOPHARMACEUTICALS GMBH.
     1772 GGAGGAGGAGGCGAGGAGGCGGC 1795
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                                      21;
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                                                       Similarity
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                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence represents a non-G-motif oligonucleotide
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                                                       0.8%;
                                                                                                        4 C; 16 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      75pp; English
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                                                                       Score 19.2;
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                                      Mismatches
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                                                                       Length 24;
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                                                                                            Synthetic.
                                                                                                          Polymerase chain reaction; mutation; mutagenesis; insertion; repetition; amplification; ss.
                                    09-AUG-1991;
                                                  09-AUG-1991;
                                                                               US5279952-A.
                                                                                                                                                14-JUL-1994
                                                                                                                                                              AAQ55605;
                                                                                                                                                                           AAQ55605 standard; DNA;
                      (TEXA)
                                                                18-JAN-1994
                                                                                                                                flanking sequence primer for manipulation
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                      VIND
                                                                                                                                                                                                                GGAGGCGGAGGCGGAGGC 24
                      TEXAS SYSTEM
                                                                                                                                                (first entry)
                                     91US-00743245
                                                  91US-00743245
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WPI; 1994-034337/04

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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic flanking sequences (see AAQ55601) were used to illustrate the novel method; a direct repeat of a specific cloned region of DNA which lies between the flanking sequences can be constructed using primers having the sequences in AAQ55602-Q55605
                                           The human and murine osteogenic protein-1 (OP-1) non-coding sequences can be used in the construction of expression vectors comprising a reporter gene which has the non-coding sequence lying contiguous to the reporter gene, the non-coding sequence being able to act on and affect expression of the reporter gene when bound to by candidate compounds. The method is used to identify compounds capable of modulating OP-1 expression. The vector may optionally comprise a second non-coding sequence and the non-coding sequence(s) used define at least one, preferably 1-6, WT1/EGR binding elements, at least one FTZ (Fushi-Tarazu) binding element or a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Construction to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Osteogenic protein; OP-1; reporter gene; screening; identification; intron; non-coding sequence; ss.
                                                                                                                                                                                                           Isolation of compounds to modulate OP-1 expression - by monitoring expression changes in a cell transformed to express osteogenic protein-1 and having additional steroid binding site.
                                                                                                                                                                                                                                                                                                                       (CREA-) CREATIVE BIOMOLECULES
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                                                                                                                                                                                                                                                                  1996-040236/04.
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                              elements, at least
binding element
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                                                                                                                                                                                 Page 58; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 7 A; 0 C; 13 G; 0 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry
                                                                                                                                                                                                                                                                                           Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TCC binding site
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Query Match

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14 C; 0

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Score 18.4; 7 T; 0 U;

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Query Match Best Local S Matches 19

l Similarity 19; Conserv

0.7%; larity 95.0%; Conservative

Score 18.4; D Pred. No. 18; 0; Mismatches

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밂 0 Other;

Length 21;

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Sequence 21

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                                     CC the expression of osteogenic protein-1 (OP-1) uses a cell transfected with a nucleic acid sequence comprising a reporter gene and an upstream concording sequence from OP-1 (OP-1) is a tissue morphogenic protein. The method is useful for screening compounds capable of stimulating or cinhibiting transcription and/or translation of the OP-1 gene, as well as compounds which may be used as therapeutics for in vivo and ex vivo correcting and alleviating a diseased condition or to regenerate lost or correcting and alleviating a diseased condition or to regenerate lost or che differentiated phenotype of cells in culture. Morphogen expression inhibiting compounds may also be used to maintain viability of the differentiated phenotype of cells in culture. Morphogen expression cinhibiting compounds identified by the new method can be used to modulate the degree and/or timing of morphogen concentration. Compounds which upcregulate levels of circulating OP-1 in vivo can be used to correct bone metabolism diseases such as osteogorosis. This sequence contained in the cupstream region of the osteogenic protein-1 (OP-1) gene. The DNA binding proteins Wt-1 and Egr-1 bind to and control transcription of DNA binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 95.0 Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-FEB-1992;
01-NOV-1993;
07-JUN-1994;
23-MAY-1995;
24-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                            Screening for compounds able to modulate osteogenic protein-1 (OP-1) expression by incubating a candidate compound with a nucleic acid wit reporter gene operatively associated with an OP-1 non-coding nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Osteogenic protein-1; OP-1; morphogenic protein; human; osteoporosis; morphogen concentration; bone metabolism disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human OP-1 Wt-1/Egr-1
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                             upstream region of proteins Wt-1 and E sequences at these
                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-422077/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oppermann H,
                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CREA-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CREATIVE
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95US-00449700.
95US-00449699.
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93US-00147023.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95US-00486343
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                sites
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
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0; Mismatches
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1772

GGAGGAGGAGGCGAGGAGG

1791

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RESULT 63
ABK99278/c
ID ABK99278/c
AC ABR9927
XX ABK9927
XX ABK9927
XX Hepati
XX Hepati
XX Hepati
XX Synthe
XX US200;
PD 30-MA'
PP 06-AP
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XX (ZHON
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                                                                                                                                                                                   The invention relates to a replicase complex comprising a hepatitis C C virus (HCV) NS5B replicase protein, a linear nucleic acid template and a CC complementary nucleic acid primer which is annealed to the 3, terminus of CC the template, where the template is at least three nucleotides and the complex is two or three nucleotides, and the template and primer do not form a stable duplex in solution in the absence of the HCV NS5B protein. The complex is useful for detecting HCV replicase activity and permits cestablishment of sensitive RNA-dependent RNA polymerase assays to screen and evaluate antiviral inhibitors and to improve the specificity and CC of a reliable system for determining kinetic and thermodynamic constants of HCV NS5B-catalysed nucleotide incorporation and investigation of CC specifically, the short RNA template and primer pairs are useful in Screening assays which are used for determining kinetic, thermodynamic and mechanistic properties of NS5B replication and ultimately in the CC development of inhibitors of NS5B replicase activity may be used for developing anti-HCV pharmaccuticals. Sequences ABK99271-ABK99296 represent HCV NS5B replicase RNA synthesis
                                                                 Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of HCV NS5B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel replicase complex comprising hepatitis C virus NS5B replicase, a 3 nucleotide-long template to which a 2 nucleotide-long primer is annealed and template and primer which do not form a stable duplex in the absence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                        Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Page 6; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zhong W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (YONG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-APR-2000; 2000US-0195852P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-APR-2001; 2001US-00828034.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2002064771-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis C virus; HCV; NS5B replicase; ss; RNA polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK99278;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK99278 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (FERR/)
21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                   l Similarity 95.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HONG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FERRARI E.
                                GGAGGAGGAGGCGGAGGAGG 1791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hong Z,
                                                                                                                                        BP;
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                                                                                                                                        A
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                                                                                0.7%;
                                                                                                                                        14 C; 0
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                                                                 Score 18.4; Di
Pred. No. 18;
0; Mismatches
                                                                   0;
                                                                                                                                      G; 0 T; 7 U; 0 Other;
                                                                                                    DB 1; Length 21;
                                                                     Indels
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RESULT 65
AAZ21462/c
ID AAZ214
XX
AC AAZ214
XX

AAZ21462 standard; DNA; 20

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AAZ21462

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1274 GGCTGAGGGCAGAGACCCTC 1293

GGCTGAGGGCAGAGACTCTC

21

Query Match Best Local S Matches 19

l Similarity 19; Conserv

0.7%; Score 18.4; larity 95.0%; Pred. No. 2: Conservative 0; Mismatche

Mismatches

DB 1; ۲.

Length 23; Indels

0

Gaps

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RESULT 64
ANQIO661
ID ANQIO
XX ANQIO
XX ANQIO
XX ANQIO
XX ANQIO
XX EST
XX EST
XX Human
XW Human
XW EP414
XX 27-FH
XX 20-AU
XX 27-FH
XX 21-JU
PF 20-AU
XX 11-SE
PR 11-
                                                          for an HLA locus, together with a second primer making up a locus-
specific primer (LSP) pair. The primers pref. define a DNA sequence that
contains all exons that encode allelic variability associated with the
HLA locus together with at least one of the adjacent intron sequences.
For Class II loci the variable exon is the second exon. The primers are
pref. located so that a substantial portion of the amplified sequence
corresponds to intron sequences. Direct deter- mination of the haplotype
is possible, providing useful information for identity of individuals for
e.g. paternity case and forensic investigations. See also AAQ10621-
Q10669. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-AUG-1989;
11-SEP-1989;
16-JAN-1990;
11-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human leukocyte antigen; major histocompatibility complex; MHC; restriction fragment length polymorphic analysis; RFLP; tissue typing; allele; PCR; ss.
      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection of adjacent and non:adjacent locus, e.g. HLA alleles amplifying genomic DNA, for direct determination of haplotype.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Simons MJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 29;
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(JEAN-)
(SIMO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-FEB-1991
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29-APR-1991
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                                                                                                                                                                                                                                                                                                                                                                                      primer is specific for nt 509-532 of HLA Class II DQwlva allele of DQB1 locus. It is used in a method for the prodn. of RFLP fragment:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) GENETYPE AG.
) GENETYPE AG.
) SIMONS M J.
      23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 49; 53pp; English.
   BP; 4 A; 7 C; 8 G; 4 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  locus-specific
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89US-00405499.
90US-00465863.
90US-00551239.
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          0 Other;
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RESULT 66
AAF73052/c
ID AAF730
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AC AAF730
XX AAF30
XX Human
XX Human
XX Antise
KW Antise
KW antiin
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Matches
                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes the use of mitotic check point genes (MCPGs) in the diagnosis and treatment of cell proliferative disorders. A method has been developed for diagnosing a cell proliferative disorder in a subject associated with a MCPG, by determining the presence of a mutant MCPG in the sample where the presence of a mutant MCPG in the sample where the presence of a mutant MCPG in the sample is indicative of a cell proliferative disorder. The method can be used for diagnosing a cell proliferative disorder such as a neoplasm, e.g. breast or colorectal neoplasm. It can also be used for treating a cell proliferative disorder, benign hyperplasia or neoplasia, particularly colon or breast cancer. It can also be used for treating disorder associated with insufficient cell proliferation or undesirable cell degeneration. The present sequence represents a PCR primer used to amplify human BUBRI, in an example from the present invention. Loss of a MCPG is associated with the mutational inactivation
Antisense oligonucleotide; daxx; inhibition; phosphorothioate; Fas binding protein; CENP-C binding protein; dap6; EAP; cytost; antiinflammatory; death associated protein 6; Ets-1 associated
                                                                                         24-APR-2001
                                                                                                                      AAF73052
                                                                                                                                                 AAF73052
                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Use of mitotic checkpoint and treatment of cell proproliferation of cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; BUB1; BUBR1; hBUB1; hBUBR1; mutation; mitosis; diag microsatellite instability; MIN; tunour; mismatch repair; chromosomal instability; detection; cell proliferative disneoplasm; breast cancer; colorectal cancer; fibrotic disobenign hyperplasia; neoplasia; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9947638-A2
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                                                                                                                                                                                                                                                                                   18; Conserv
                                                                                                                                                                                                                                                                                                                                                                          human BUB1 gene
                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                 standard; DNA;
                                                                                                                                                                                                                                                       CCAGAAGCAGCTGCGTTTT 1391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 34; 56pp; English
                                                        inhibitory antisense phosphorothicate oligonuclectide
                                                                                                                                                                                                                                                                                                                                               BP; 5 A; 5 C; 6
                                                                                                                                                                                                                                                                                     Conservative
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                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US005692
                                                                                        entry)
                                                                                                                                                                                                                                                                                                  94.7%;
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                                                                                                                                                                                                                                                                                                                                               G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                           ...
                                                                                                                                                                                                                                                                                                   Score 17.4;
Pred. No. 2
                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                developing methods for the diagnosis disorders or for increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lengaver
                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                                                                                                                                                               Length 20;
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protein;
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AAC925

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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes an antisense compound (I) up to 30 nucleobases in length, where (I) inhibits expression of daxx (also known as Fas binding protein, CRP-C binding protein, daps for death associated protein 6 and EAP for Ets-1 associated protein). (I) has cytostatic and antiinflammatory activity, and can be used in antisense therapy and as a modulator of daxx. (I) is useful for inhibiting the expression of daxx in cells or tissues in vitro. (I) can be utilised for diagnostics, therappeutics for the treatment of diseases associated with the expression of daxx, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation and as research reagent. The present sequence represents an inhibitory human daxx antisense phosphorothicate oligonucleotide which is used in the avantisense phosphorothicate oligonucleotide which
                                                                                                                                                                                                                                                                                   ribosome transport; cytokinėsis; nūcleogenesis; cell cell growth; transcriptional repression; replication; signal transduction; chromatin decondensation; Ag-NOR nucleolin antibody; systemic connective tissue diseas
                                                                                                                                                                                                                                                                                                                                                                      Human nucleolin; P92; C23; phosphoprotein; ribosome biogenesis; ribosome transport; cytokinesis; nucleogenesis; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human nucleolin phosphorothicate antisense oligonucleotide,
                                                                                                                                                                                                  scleroderma-like chronic graft versus host expression inhibition; tumour formation; ca
                                                                                                                                                                                                                                                          systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC92593;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC92593 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antisense compounds capable of modulating expression of daxx useful for diagnosis, prophylaxis and treatment of diseases associated with expression of daxx.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Col 49; 59pp; English.
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Pred. No. 26
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disease; SLE;
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RESULT
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ribosomes in the nucleolus via a ribonucleoprotein consensus sequence. However, it has also been shown to be involved in cytokinesis, nucleogenesis, cell proliferation and growth, transcriptional repression, replication, signal transduction, and chromatin decondensation. Nucleolin is a member of the Ag-NOR (active ribosomal gene located in the nucleolar organiser region) family of proteins which are markers of active ribosomal genes, and whose expression is associated with the prediction of tumour growth rate. The presence of antibodies against nucleolin are associated with systemic connective tissue diseases such as systemic lupus erythematosus (SIB) and scleroderma-like chronic graft versus host disease. The oligonucleotides of the invention are useful for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotides were designed to target different regions of the human nucleolin mRNA, and were analysed for their effect on nucleolin mRNA levels by quantitative real-time PCR. Nucleolin (also known as P92 or C23) is the most abundant nucleolar phosphoprotein in actively growing cells. Nucleolin primarily participates in ribosome biogenesis and transport of ribosomal components, being able to transiently bind to pre-
                                                                                                                      Candida albicans Synthetic.
18-FEB-2000; 2000US-0183534P
                              20-FEB-2001; 2001WO-US005551
                                                                                                                                                                                                              Primer A #30
                                                                                                                                                                                                                                            04-DEC-2001
                                                                                                                                                                                                                                                                                                       AAS23716 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel antisense compound targeted to human nucleolin which specifically hybridizes with and inhibits the expression of human nucleolin, useful for modulating the expression of nucleolin in cells.
                                                                                         WO200160975-A2
                                                                                                                                                                                                                                                                          AAS23716;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prevention and treatment of conditions associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences AAC92560-C92639 represent antisense oligonucleotides targetted to the human nucleolin gene, which inhibit its expression. The antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; Col 41-42; 41pp; English.
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                                                                                                                                                                identification;
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94.7%;
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Z
                                                                                                                                                             essential gene; GRACE; pathogenic fungus; conditional expression; fungal infection;
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Pred. No. 2
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0; Mismatches
                                                                                                                                                                                                         identifying C. albicans GRACE
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in the nucleolar
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RESULT 69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to novel methods for constructing fungal strains useful for identification and validation of gene products as targets for therapeutic agents, for creating a collection of identified essential genes, and screening assays for the discovery of new drugs. The invention provides the GRACE (gene replacement and conditional expression) method for the construction of mutant organisms referred to as GRACE strains of the organism. The invention can be applied to any organism, particularly a pathogenic fungus e.g. Candida albicans, Asparallus funigatus and Cryptococcus neoformans. The methods are useful to identify agents that may be used in the treatment of fungal infections. AAS23687-AAS23747 represent primers A #1-61 used as probes
                                                                                                                                              29-DEC-2000; 2000US-0259128P
20-FEB-2001; 2001US-00792024
22-AUG-2001; 2001US-0314050P
                                                                                                                                                                                                                                                                                               Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                                                                                                                                                 26-DEC-2001; 2001WO-US049486
                                                                                                                                                                                                                                                                           Candida albicans
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                                                                                                                                                                                                                                                  WO200253728-A2.
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                                                                                                                       (ELIT-) ELITRA PHARM INC.
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                                                                        2002-566694/60.
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18; Conserv
                                                                                                                                                                                                                                                                                                                                                    albicans GRACE strain PCR
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94.7%;
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Claim 36; SEQ

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4054; 167pp +

Sequence Listing; English

expression.

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Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of the gene and placing other allele of the gene under conditional expression

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ADDULT 70
ADD71322
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Best Local S
Matches 18
                                                                       Detecting the presence of a target nucleic acid molecule in templates combining a detection probe, a first oligonucleotide, second oligonucleotide, a primer and templates suspected of containing the target nucleic acid molecule.
                                                                                                                                                                                                                                                                             WPI; 2003-803888/75
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nilarity 94.7%;
Conservative
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signal transduction, SH2 domain, SH3 domain, src homology integrin signalling, receptor tyrosine kinase signalling; growth factor receptor signalling; PINCH; v-Abl; Ras; Sos; transcriptional activation; cancer; tumour; leukaemia; breatings; b Homo sapiens expression Nck-2; adapter protein; Nck adapter protein; hNck-beta; Grb4; ltransduction; SH2 domain; SH3 domain; src homology domain; antisense leukaemia; breast oligonucleotide;

US6165728-A

19-NOV-1999; 99US-00444053.

19-NOV-1999; 99US-00444053.

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2001-090480/10

Novel antisense compound which inhibits expression for treating disease or condition associated with e and as research reagents, kits and diagnostics. n of human nck-2 useful expression of nck-2,

Claim 1; Col 41-42; 38pp; English.

Sequences AAC92649-C92728 represent antisense oligonucleotides targetted to the human Nck-2 gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human Nck-2 mRNA, and were analysed for their effect on Nck-2 mRNA levels by quantitative real-time PCR. Nck-2 (also known as Nck adapter protein, hNck-beta and Grb4), contains both SH2 and SH3 src homology domains and functions as an adapter protein in integrin-mediated and receptor tyrosine kinase-mediated signal transduction, particularly in growth factor receptor signalling Moreover, Nck-2 participates in pathways that connect growth factor receptor signalling moreover, Nck-2 participates in pathways that that a its

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RESULT 72
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                          The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; bypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                            Claim 15;
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RESULT 73
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Best Local S
Matches 18
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Best Local S
Matches 18
                                                                                                                  The present invention relates to coding sequences (ADC49210 and for rabbit hyaliuronic acid synthetase (HAS)-2 or HAS-3 (ADC4921 ADC49213). The sequences of the invention can be used for treat joint disorders and articular diseases, such as osteoarthritis. present sequence was used to illustrate the invention.
                                                                                                                                                                                                                    acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fire, wipo.int/pub/published_pct_sequences
                                                                                            Sequence
                                                                                                                                                                                                                                                         WPI; 2003-508645/48.
                                                                                                                                                                                                                                                                                                      27-JUL-2001; 2001JP-00228543
                                                                                                                                                                                                                                                                                                                                                                                 JP2003038185-A
                                                                                                                                                                                                                                                                                                                                                                                                                                articular disease; osteoarthritis; HAS-1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Rabbit; hyaluronic acid synthetase; enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC49221;
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                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
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                      2033
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                                                                                                                                                                                                                  l gene useful for nucleic acid sequencing, code synthetase and has hyaluronic acid synthetase
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nilarity 90.0%;
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Pred. No. 3
                                                       Score 16.8;
Pred. No. 34;
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                                                                                                                                                                                                                                codes rabbit hyaluronic
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                               This is the nucleotide sequence of primer Exon 2n, which was used with primer Exon 1 (see AAV64912) in RT-PCR to characterise the splice junctions of the latency associated transcript (LAT) of herpes simplex virus type 1 (see AAV64883-86). The invention relates to methods of stabilising unstable gene transcripts. A claimed polynucleotide comprises: a polynucleotide encoding a gene product; a 5'-sequence of an intron, including the splice donor and splice acceptor sites (see AAV64885-86), and a 3'-sequence of the same intron, including a hairpin structure (see AAV64887) next to the intron's branchpoint. A preferred intron is the 2.0 kb LAT of a herpes virus. Methods and compositions using the polynucleotide can be used in gene therapy and more generally as research reagents, markers of gene production, in therapeutic or diagnostic compositions, in drug screening and to identify transcripts produced only at selected stages of the cell cycle
                         RT-PCR primer used to amplify CHYM cDNA.
                                                                                  ABQ78582
                                                                                                         ABQ78582 standard; DNA; 18
                                                                                                                                                                                                                                                                                         Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-609982/51
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Pred. No. 39;
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Human; mucin 4; MUC4; peripheral blood monocyte; PBMC;

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RESULT 76
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Best Local :
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                                                                                                                                                                                                                                                                                                                                          Human; casein kinase 2-alpha prime; diabetes mellitus; hyperproliferative disorder; breast cancer; prostate cancer; liver cancer; inflammation; tumour formation; cytantidiabetic; antiinflammatory; antimicrobial; phosphorothio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human casein kinase 2-alpha prime antisense oligonucleotide #14.
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                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                 antisense therapy; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-DEC-2000; 2000US-00733444
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Pred. No. 31;
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ABT4340/c
ID ABT434
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AC ABT434
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DT 22-SEP
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PF 30-MA)
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PR 24-AUC
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XX WPI;
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Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to antisense oligonucleotides and methods for modulating the expression of human or mouse casein kinase 2-alpha prime. The antisense oligonucleotides are useful for inhibiting the expression of casein kinase 2-alpha prime, and for treating diseases or conditions associated with aberrant expression of casein kinase 2-alpha prime. Such diseases include diabetes mellitus, and hyperproliferative disorders (particularly cancers e.g. breast cancer, prostate cancer, or liver cancer). The antisense compounds are also useful for diagnostics, inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. ABS67840-ABS67917 represent human or mouse casein kinase 2-alpha prime antisense oligonucleotides which comprise a phosphorothicate
                                                                                                                          31-MAY-2001;
24-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides targeted to nucleic acid encoding casein kinase 2-alpha prime, useful for diagnosing and/or treating a disease or condition associated with expression of casein kinase 2-alpha prime.
  Nucleic acids isolated
                              WPI; 2003-167523/16.
                                                                                                                                                                 30-MAY-2002; 2002WO-JP005295
                                                                                                                                                                                                                         WO2002103017-A1
                                                                                                                                                                                                                                                                              Neuroblastoma;
                                                                                                                                                                                                                                                                                                          Neuroblastoma-related DNA
                                                                                                                                                                                                                                                                                                                                                                 ABT43400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                                                                                                                                                                                                   22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                           ABT43400 standard; DNA; 20
                                                                                                 (CHIB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-FEB-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                      77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1881 CTGGAGGAGGAGGAG 1898
                                                                                   CHIBA PREFECTURE
HISAMITSU PHARM (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SISI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                           2001JP-00163666.
2001JP-00255260.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM INC
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                                                                                                                                                                                                                                                                                                                                   (first entry
                                                                                                                                                                                                                                                                              prognosis;
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 from neuroblastoma showing
                                                                                   COLTD
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                                                                                                                                                                                                                                                                                                         sequence #315.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred.
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                                                                                                                                                                                                                                                                              oligonucleotide
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1545 ACCTGGCTGAACATTGCA 1562

Query Match Best Local S Matches 17

17; Conserv

Conservative

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Score 16.4; D. Pred. No. 41; 0; Mismatches

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RESULT 78
ABT32545/c
ID 32545/c
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Best Local S
Matches 17
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ABT32545
                                                The invention comprises nucleic acids that show increased expression in human neuroblastomas with poor prognosis over those with a good prognosis. The nucleic acids of the invention are useful as a tool for distinguishing neuroblastomas with a favourable prognosis (spontaneous regression) from neuroblastomas with a poor prognosis (high malignancy) The DNA sequences ABT32224 - ABT32571 represent oligonucleotides used in an example of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human neuroblastoma with good prognosis, prognosis of neuroblastoma and providing
                                                                                                                                                                                                                                                                          Example
                                                                                                                                                                                                                                                                                                                        Nucleic acids having higher expression prognosis for diagnostic prediction of
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-140476/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHIB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAY-2001;
24-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200297093-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    high malignancy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Neuroblastoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABT32545;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention comprises DNA sequences that show enhanced expression in human neuroblastoma with good prognosis. The bNA sequences of the invention are useful in clarifying good/poor prognosis of neuroblastoma. The present DNA sequence was used in the exemplification of the inventic
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nakagawara A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neuroblastoma-related oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 25(1); 444pp; Japanese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHIBA PREFECTURE
HISAMITSU PHARM (
                                                                                                                                                                                                                                                                    5; Page 28; 111pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA;
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2001JP-00255226.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prognosis; spontaneous regression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
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Pred. No. 41;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                           in human neuroblastoma with poor neuroblastoma prognosis.
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genetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              probe; ds;
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RESULT 79 ABZ81534/c ID ABZ81534 :

standard;

DNA;

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Best Local S
Matches 17
                                                                                                                                        The present invention relates to novel antisense oligonucleotides (ABZ81522-ABZ81593) which are targeted to human protein kinase A (PKA) regulatory subunit RII beta nucleotide sequence (ABZ8113), and which specifically hybridise with and inhibit the expression of the PKA regulatory subunit RII beta (PKA is also known as CAMF-dependent protein kinase). The antisense oligonucleotides are useful for modulating the expression of PKA regulatory subunit RII beta and for treating diseases or conditions associated with aberrant expression of PKA regulatory subunit RII beta, e.g. diabetes or cancer. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway
                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                  Seguence
                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides targeted to nucleic acid encoding protein kinase A regulatory subunit RII beta, useful in treating diseases e.g. cancer associated with the aberrant expression of the protein kinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-239434/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JUL-2002; 2002WO-US022629.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003010283-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PKA regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       regulatory subunit cancer; infection;
                             1770 GAGGAGGAGGCGGAG 1787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B₽,
                                                                                                                                                                                                                                                                                                                                                3; Page 74; 98pp; English.
 20
                                                                       Similarity
                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wyatt JR
                                                                                                                 BP; 0
                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             subunit RII beta antisense oligonucleotide ISIS #114459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "Oligonucleotide has phosphorothicate backbone
all cytidine nucleotides are 5-methylcytidine. Optiona
some nucleotides with 2'-methoxyethyl (2'-MOE wings)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modification"
                                                                                                                 A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antidiabetic; antisense therapy; phosphorothioate; bitor; protein kinase A; PKA; RII beta; cAMP-dependent protein kinase; diabetes; inflammation; tumour; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                     0.6%;
                                                                                                                 12 C; 4 G; 4 T;
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                                                       Score 16.4; I
Pred. No. 41;
0; Mismatches
                                                                                                                 0
U;
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                                                                                                                 0 Other;
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                                                                                   Length
                                                         Indels
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                                                       Gaps
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Antisense gene

therapy; RAIDD; death domain;

caspase

recruitment

Mouse RAIDD antisense oligonucleotide #75.

21-OCT-2002

(first

entry)

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RESULT 80
AAF73054/c
ID AAF730
XX AAF730
XX AAF730
XX AAT188
AC Human
XX Ant188
KW Fas bi
KW Fas bi
KW Fas bi
KW Ant116
KW ANT186
KW Fas bi
KW ANT186
KW Fas bi
KW ANT186
XX US6180
XX INSTENDING
PF 24-JAN
XX US6180
XX
RESULT E
ABK9982]
ID ABH
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an antisense compound (I) up to 30 nucleobases in length, where (I) inhibits expression of daxx (also known as Fas binding protein, CRNP-C binding protein, daps for death associated protein 6 and EAP for Ets-1 associated protein). (I) has cytostatic and antiinflammatory activity, and can be used in antisense therapy and as a modulator of daxx. (I) is useful for inhibiting the expression of daxx in cells or tissues in vitro. (I) can be utilised for diagnostics, therapeutice for the treatment of diseases associated with the expression of daxx, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation and as research reagent. The present sequence represents an inhibitory human daxx antisense phosphorothioate oligonucleotide which
                                                                                                                                                                    ABK99821;
                                                                                                                                                                                                                        ABK99821 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense oligonucleotide; daxx; inhibition; phosphorothioate; Fas binding protein; CENP-C binding protein; dap6; EAP; cytostatic; antiinflammatory; death associated protein 6; Ets-1 associated protinfection; inflammation; tumour formation; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JAN-2000; 2000US-00490692
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule (II) encoding RAIDD which is an adaptor molecule containing both death domain (DD) and caspase recruitment domains (CARD), where (I) specifically hybridises with and inhibits expression of RAIDD, or specifically hybridises with and 8-nucleobase portion of an active site on (II). (I) is useful for conhibiting the expression of RAIDD (Receptor interacting protein (RIP) associated ICH-1/CED-3-homologous protein with death domain) in cells or tissues, and for treating an animal having a disease or condition associated with RAIDD, where the disease or condition is a hyperproliferative disorder such as conditions of growth or metabolic disorder. (I) is also useful for diagnostics, therapeutics, prophylaxis, as research reagents and kits, for distinguishing functions of various members of a biological pathway, and in antisense gene therapy. (I) is also useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation. This sequence represents a mouse RAIDD cantisense oligonucleotide used to control expression of the RAIDD protein
                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 16
                                Antisense gene therapy; RAIDD; death domain; caspase recruitment domain; CARD; hyperproliferative disorder; cancer; growth disorder; mouse; metabolic disorder; infection; inflammation; tumour formation; RIP associated ICH-1/CED-3-homologous protein with death domain;
                                                                                                                       Mouse RAIDD antisense oligonucleotide #74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antisense compound that hybridizes and inhibits nucleic acid encoding RAIDD which is an adaptor molecule containing both death and caspase recruitment domains, for treating hyperproliferative
                 receptor interacting protein; antisense oligonucleotide;
                                                                                                                                                                                                                                   ABK99820 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 4
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                                                                                                                                                                                                                                                                                                                                                            1227
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                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                          CTCCAGCATGTGCTGG
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                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                             entry)
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Pred. No.
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The invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule (II) encoding RAIDD which is an CC adaptor molecule containing both death domain (DD) and caspase contributed to an active site of (I) specifically hybridises with and CC inhibits expression of RAIDD, or specifically hybridises with at least an CC 8-nucleobase portion of an active site on (II), (I) is useful for CC inhibiting the expression of RAIDD (Receptor interacting protein (RIP) CC associated ICH-1/CED-3-homologous protein with death domain) in cells or tissues, and for treating an animal having a disease or condition CC theorems, (I) is also useful for diagnostics, therapeutics, prophylaxis, as research reagents and kits, for distinguishing functions of various members of a biological pathway, and in antisense gene therapy. (I) is also useful prophylaxis, and in antisense gene therapy. (I) is also useful prophylaxis, and in antisense gene therapy. (I) is also useful prophylaxis, and in antisense gene therapy. (I) is also useful prophylaxis, and in antisense gene therapy. (I) is also useful prophylaxis, and in antisense gene therapy. (I) is also useful prophylaxis, and in sequence represents a mouse RAIDD continuous of the CC antisense oligonucleotide used to control expression of the RAIDD protein
Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel antisense compound that hybridizes and inhibits nucleic acid encoding RAIDD which is an adaptor molecule containing both death and caspase recruitment domains, for treating hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zhang
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                                                                                                                                                                                                                                                                                                                                                                                                                               English.
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Other;
                                      RAIDD protein
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RESULT 83
AAQ92210/
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AC AAQ92
XX 12-JJ
XX 12-JJ
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KW Prime
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AAQ92210
10-NOV-1994;
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                                      WO9513397-A1
                                                            Synthetic
                                                                               benign; malignant; neoplasm; lung;
                                                                                                  Primer; polymerase chain reaction; amplify; mutant; K-ras;
                                                                                                                        p53 detection
                                                                                                                                            12-JAN-1996
                                                                                                                                                                 AAQ92210;
                                                                                         flanking
                                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                          Similarity
                                                                               region; amplification; probe; detection; sputum; maliqnant; neoplasm; lung; lung cancer; head; neo
                                                                                                                                                                                    standard; DNA; 17
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                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                            (first
                                                                                                                       probe,
94WO-US012947
                                                                                                                                            entry)
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Pred. No.
                                                                                                                                                                                                                                                                                Mismatches
                                                                              lung cancer;
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RESULT 84
AAF02815/C
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XX Ribo;
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequences given in AAQ92112-211 are probes which were used in the detection of a mutant p53 gene sequence. The DNA to be detected is amplified using PCR and then these probes which are pref. labeled using 32-P gamma-ATP are used to detect the mutant sequences. The primers and probes given in AAQ92098-219 are used in the method of the invention for detecting mammalian target DNA in sputum samples. Analysis of the target DNA is used to diagnose benign or malignant neoplasms of the lung. It is also useful for screening people at high risk or for monitoring progress of treatment of lung neoplasms. The method is based on the discovery that mutant target DNA associated with lung cancer is present at detectable levels in sputum. Cells shed into sputum from head and neck cancers may also be detected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting
diagnosis
or p53 tum
                                                  useful for interferon
                                                                    Enzymatic and antisense nucleic acid inhibition of repressor useful for producing e.g.\ granulocyte colony stimulating fact
                                                                                                                                                    WPI; 2000-647423/62
                                                                                                                                                                                                                                                                                                                                                11-APR-2000;
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                                                                                                                                                                                                                                                                                                12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               interferon alpha;
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                                               producing
                                                                                                                                                                                                    Zwick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 36; 122pp;
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                                                                                                                                                                                                    Pavco
                                             e.g. granulocyte erythropoietin.
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94.1%;
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Claim

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Claim 88;

Page 131; 200pp; English

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RESCUENT AND PROCESS OF PROCESS O
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.2359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; NAZyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHI; lymphocytic leukaemia; human immunocytoma; IWC; immune thrombocytopaenia; stroke; dementia; MCL; immunocytoma; IWC; immunocytoma; IWC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheiner's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; cerebrovascular accident; CVA; Alzheiner's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interest the production of the production of genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                               Nucleic acid molecules, e.g., enzymatic nucleic constructs, which down regulate expression of a growth inhibitor gene useful for treating, e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                           central nervous system injury.
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(BLAT/)
(MCSW/)
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BLATT L.
MCSWIGGEN J.
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16; Conser
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nilarity 94.1%;
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Pred. No. 4:
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RESULT 86
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Best Local S
Matches 16
26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-CCT-2000; 2000GB-00024263.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLF muscle; myosin; chromosome 22; gene therapy; vaccine; heart skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                    25-MAY-2001;
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16; Conservative
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The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-CC 1 can be used in gene therapy and vaccine production. The hGDMLP-1 CC mucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 CC protein variants having desired phenotypic improvements, and for CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 (c) proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapy and for surface-enhanced laser desorption ionisation, as the propential sequence sencoding hGDMLP-1 may be used for diagnosing a CC objunction, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a CC objunction and in vaccines of the present therapy. The propens to sequence represents an oligomer used for diagnosing of the present sequence in the expression of hGDMLP-1 in particular heart CC in proteins, but was obtained in electronic format directly from WIPO as the printed of the printed on the present invention. N.B.
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30-JAN-2001;
05-FEB-2001;
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Sequence 17 BP; 6 A; 2 C; 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 929; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polypeptide, or as specific b
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                                    ftp.wipo.int/pub/published_pct
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2001WO-US000666.
2001WO-US000667.
2001WO-US000668.
2001WO-US000669.
2001WO-US000669.
2001WO-US000670.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e, for raising antibodies that recognize hGDMLP-1 proteins, biomolecule capture probes for surface-enhanced laser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hanzel
<u>ი</u>
2 T; 0 U;
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                                      gequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   뮸
0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chen
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            S
                           Query Match
Best Local Similarity
Matches 16; Conserv
       1264 AGCTGGAAGAGGCTGAG 1280
μ
                            Conservative
                                  0.6%;
17
                           0;
                                  Score 15.4;
Pred. No. 42;
                             Mismatches
                                          B
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                            1;
                                         Length 17;
                            Indels
                            0
                            Gaps
                             0
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RESULT 87
ABNO8667
ID ABNO8
XX ABNO8
XX ABNO8
XX ABNO8
XX 29-MA
XX Human
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; genome-derived myosin-like protein 1; GDMLP-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human GDMLP-1 17-mer scanning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABN08667;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               muscle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABN08667 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       myosin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosome 22; gene therapy; vacc disorder; amplicon; screening; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:5 sequence SEQ ID NO:8659.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rotein 1; GDMLP-1; hGDMLP-1; heart;
therapy; vaccine; heart disease;
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WO200192524-A2

skeletal muscle

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RESULT 88
ABV78925
ID ABV78
XX
AC ABV78
XX
DT 03-JJ
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                                                                                                                                                                                                                                                                                                                   CC The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-CC 1 can be used in gene therapy and vaccine production. The hGDMLP-1 CC nucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 CC protein variants having desired phenotypic improvements, and for CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognize hGDMLP cc used as immunogens to raise antibodies that specific biomolecule cc proteins, as standards in assays used to determine the concentration cc and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therape training the protein, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a cdisorder associated with the expression of hGDMLP-1, in particular heart cc and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

CC The present sequence represents an oligomer used in the screening of the present sequence data for this patent did not form part of the printed capture with mile and not form part of the printed controlled and the printed on the protein directly from WIPO at the wipo int/ml/humilised for the present directly from WIPO at the purpose.
                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 16
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
                  03-JAN-2003
                                                    ABV78925
                                                                                   ABV78925
                                                                                                                                                                                                                                                                                                                   Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser description ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                          1529 GCTGGAGGAGGCCAAGA 1545
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                                                                                                                                                                                                                                              l Similarity
16; Conserv
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                                                                                     standard;
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                                                                                                                                                                                                                                                Conservative
                (first entry)
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2000US-0234687P
2000US-0236359P
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2001WO-US000662.
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                                                                                                                                                                                                                                                                                                                 A; 3 C; 7
                                                                                     DNA; 17
                                                                                                                                                                                                                                                             0.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanzel
                                                                                                                                                                         17
                                                                                                                                                                                                                                                                                                                 G; 1
                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                            Score 15.4; I
Pred. No. 42;
0; Mismatches
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                                                                                                                                                                                                                                                                            Length 17;
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30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001US-00864761.
                                                                                                                                                                                                                                                  Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                               Example 2; Page 86; 718pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EP1229046-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human HTPL scanning oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                        AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002EP-00001167
                                                                                                                                                                                                                                                                                                                                                            2001US-0327898P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ID 171.
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RESULT 89
ABK17920/
ID ABK1
XX
AC ABK1
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC The present invention relates to human testis expressed Patched like CC protein (HTPL, see ABVB7575 to AB79762 and ABB98519 to ABB98520). HTPL CABB Potents, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop CC codon in HTPL-S (S for short) compared to HTPL-I (L for long). HTPL CS shares an overall structure organisation with the Patched protein. The start of that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was CC mapped to human chromosome 10p12.1. HTPL and its coding sequence are cuseful for diagnosing a disorder caused by mutation in HTPL, and in CC therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human CHTPL. Such disorders include disorders of testis, or adrenal, adult and CC foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, shelter useful diagnostic markers and potential therapeutic acids are clinically useful diagnostic markers and potential therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an certample from the invention
                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 16
                                                                                         7920/c
ABK17920 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                   2120 CCACGGGGCCGCAGTGG 2136
                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 7 C; 7
                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                            RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                             0.6%;
                                                                                                                                                                                                                                             17
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Pred. No. 42;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 1 T; 0
                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
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ABK17920;

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09-APR-2002

(first

entry)

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В
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                                                                                                                                                                                                                                                                       Weber syndrome, Kippel-Trenamunay-Weber syndrome, Osler-Reber-rendu CC syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies cunder conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or constanting the cell with (I). (I) is useful for cleaving RNA of CC ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnosis of conditions and CC diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically CC argeting genes that share homology with ERG gene or ERG fusion genes. ABK17354-ABK27719 represent nucleic acide, including antisense and celsarded PCR primers of the invention
                                                                                                           Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca yulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnorary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel polynucleotide which down regulates expression of Ets-related geruseful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 69; 149pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-)
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                                               1880
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                                                                                                           l Similarity
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  GCTGGAGGAGGACGCGG 1
                                                     GCTGGAGGAGGACGAGG 1896
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                                                                                                                                                                                                                          BP; 1
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                                                                                                                                   0.6%;
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                                                                                                                                         Pred.
                                                                                                                                                                       Score 15.4;
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                                                                                                              Mismatches
                                                                                                                                                                                                                          ;
                                                                                                                                      No.
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                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                             0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mclaughlin
                                                                                                              1.
                                                                                                                                                                    Length 17;
                                                                                                              Indels
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                                                                                                              Gaps
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RESULT 90
ADB98963;

XX
ADB98963;

XX
ADB98963;

XX
AC
DEC-2003 (first e
AC
ADB98963;

XX
DE
LRP5 mutagenic PCR p
XX
DE
LRP5 mutagenic PCR
DE
LRP6 mutants, which
CC
Call. The HBM-like
pT
CC LRP6 mutants, which
CC cell. The HBM-like
DC lare1 modulation. Th
phemotype in a subje
CC Detelmodulation. Th
phemotype in a subje
CC Detelmodulation. Th
phemotype in a subje
CC cell. The PF
XX
Sequence 17 BP; 3 A;
ARAGO296
ID ARAGO
XX ARAGO
XX ARAGO
XX ID-A
XX Prim
XX Alph
KW anti
KW ss.
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                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to High Bone Mass (HBM), LRP5 (Zmax1) and LRP6 mutants, which results in a HBM-like phenotype when expressed in a cell. The HBM-like phenotype results in bone mass modulation and/or lipid level modulation. The invention is useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 6 C; 6 G; 2 T; 0 U; 0 Other.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-0291311S-
01-FEB-2002; 2002US-0353568P.
04-MAR-2002; 2002US-0361293P.
              alpha-2-delta-D; calcium channel; 12p13.3; gabapentin; cytostatic; anticonvulsant; antimigrane; antiparkinsonian; antidepressant; pri
                                                                                                                               AAA09296;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid comprising a mutation in LRP5 or LRP6, useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject
                                                               Primer
                                                                                                 10-AUG-2000
                                                                                                                                                                 AAA09296 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-MAY-2002; 2002WO-US014877
                                                                                                                                                                                                                                                                                 1349 CTTTCCCAGGGCAGCTG 1365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mass modulation; osteoporosis; PCR; primer; ss
                                                                                                                                                                                                                                                   17
                                                                                                                                                                                                                                                                                                               16;
                                                               for
                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENOME THERAPEUTICS CORP. WYETH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       from e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anisowicz
                                                              human
                                                                                                                                                                                                                                                                                                                 0.6%;
ilarity 94.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene therapy; High Bone Mass; HBM; LRPS; Zmax1;
                                                                                                 (first entry)
                                                              alpha-2-delta-D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     53; 629pp; English.
                                                                                                                                                                 CDNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Graham JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 #82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                 ΒP
                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                 Score 15.4;
                                                                                                                                                                                                                                                                                                                                     Pred.
                                                                                                                                                                                                                                                                                                                   Mismatches
                                                               gene
                                                                                                                                                                                                                                                                                                                                     No.
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                                                                                                                                                                                                                                                                                                                     Gaps
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RESULT 92
ADE30113/c
ID ADE30113;
XX
AC ADE30113;
XX
Mitogen activated protein kir
XX
DE Mitogen activated protein kir
XX
E Mitogen activated protein kir
XX
DE Mitogen activated protein kir
XX
E Mitogen activated protein kir
XX
Short interfering nucleic aci
XX
Sportstic; anorectic; antidi
XX
Inflammatory disease; asthma
XX
Sportiasis; inflammatory bowe.
XX
Synthetic.
XX
PD 04-SEP-2003.
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-OCT-1998;
30-OCT-1998;
29-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The alpha-2-delta-D gene encodes a calcium channel subunit polypeptide. The gene has been mapped to chromosome 12p13.1. This gene and the related alpha-2-delta-C and -B genes are useful for protecting mammalian cells from abnormal calcium flux by introducing expression vectors containing the respective gene into mammalian cells. The antisense genes are also useful for treating or preventing epilepsy. The alpha-delta-2-A protein is a high-affinity binding target of the anti-convulsant drug gabapentin. Therefore, alpha-delta-2 proteins may also be targeted to treat seizure-related syndromes, migraine, attaxia, vestibular defects, chronic pain, sleep interference, anxiety, amyotropic lateral sclerosis (ALS), multiple sclerosis, mania, tremor, parkinsonism, substance abuse or addiction evadrones, mood. depression or cancer
                                                                                                                                                                         short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anorectic; antidiabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antirheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human nucleic acids encoding the alpha2delta-C and alpha2delta-D proteins, useful in the treatment of epilepsy, migraine, chronic pair anxiety, multiple sclerosis or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200020450-A2
                                                                                                                                         inflammatory disease; asthma; septic shock; rheumatoid
psoriasis; inflammatory bowel disease; drug screening;
                                                                                                                                                                                                                                                                                             Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:735.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-303744/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1075 TGAGGAAGCGGTTCATG 1091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ω
                                                                                                                       engineering; pharmacogenomic; gene mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGAGGAAGCGGATCATG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 76; 88pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mood,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-0103322P.
98US-0106473P.
98US-0114088P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               depression or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 2 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8
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                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Į,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15.4;
No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      В
                                                                                                                                                           rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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ADE30322
ID ADE30
XX ADE30
AC ADE30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a short interfering nucleic acid (sina) control that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or cells trissue explants or cells trissue explants or cells translation; and (4) corresponding these vectors that express sina and cells containing these vectors. MAPK sinase cells artiarthritic, anorectic, antidiabetic, antiinflammatory, antiasthmatic, immunosuppressive, antibacterial, antirheumatic, cells, cells and early can be used to modulate the expression of MAPK genes, in cells, cells can be used to modulate the expression of MAPK genes, in cells, cells and II; a wide range of tumours, and inflammatory diseases (asthmatic septic shock, rheumatoid arthritis, psoriasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target celested; the control of maphing (e.g. of single-nucleotide control of the present sequence represents a MAPK sina which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
15-JAN-2003; 2003US-0440129P.
                      short interfering nucleic acid; siNA; downregulation; inhibition mitogen-activated protein kinase; MAP kinase; MAPK; RNA interfer cycostatic; anorectic; antidiabetic; antiinflammatory; antiasthm immunosuppressive; antibacterial; antirheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritic; psoriasis; inflammatory bowel disease; drug screening; psoriasis; inflammatory bowel disease; drug screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-JAN-2003;
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                                                                                                                                                                                                                                                                                               Mitogen activated protein kinase siNA oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE30322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
hes 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ū
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,,</u>
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Pred.
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                                                                                                                                                                                                                                             inhibition;
                                                                                                                                                                                 A interference;
antiasthmatic;
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RESULT 94
AAF52822
ID AAF52
XX AAF52
XX AAF52
XX IGF-1
XX AAF62
XX IGF-1
XX AAF62
XX IGF-1
XX IGF-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                          CC The present invention describes a short interfering nucleic acid (siNA)
CC that downregulates expression of a mitogen-activated protein kinase
CC (MAPK) genes by RNA interference. Also described: (1) a method for
modulating expression of MAPK genes in cells, tissue explants or
CC organisms by introduction of siNA; (2) kits for in vitro or in vivo
CC delivery of siNA; (3) conjugates and/or complexes of siNA; and (4)
CC vectors that express siNA and cells containing these vectors. MAPK siNAs
CC have cytostatic, annoractic, antidiabetic, antifilammatory,
antiasthmatic, immunosuppressive, antibacterial, antirheumatic,
CC antiasthmatic, immunosuppressive, antibacterial, antirheumatic,
CC antiarthritic, antipsoriatic and gastrointestinal activities. The MAPK
CC siNAs can be used to modulate the expression of MAPK genes, in cells,
CC tissue explants or organisms, e.g. for treating obesity; diabetes types I
CC tissue explants or organisms, e.g. for treating obesity; diabetes types I
CC disease). They can also be used for drug screening; diagnostis; target
CC disease). They can also be used for drug screening; diagnostis; target
CC disease). They can also be used for drug screening; pharmacogenomics;
constitution and validation, genetic engineering; pharmacogenomics;
constitution and validation, genetic engineering pharmacogenomics;
constitutions. The present sequence represents a MAPK siNA which is used
CC in the exemplification of the present invention.
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Best Local Sim.
Matches 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
29-SEP-2002;
09-SEP-2002;
15-JAN-2003;
Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris;
                                                                                                                     30-MAR-2001
                                                                                                                                                      AAF52822;
                                                                                                                                                                                         AAF52822 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein kinase genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen
                                                                                  IGF-I oligonucleotide #3782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
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                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                          CAUGGUCAUCUGUAAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ
                                                                                                                                                                                                                                                                                                                                                                                                              BP; 4 A; 4 C; 4 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; 2002US-0358580P.
2002US-0363124P.
; 2002US-0386782P.
; 2002US-0406784P.
; 2002US-0408378P.
; 2002US-0409239P.
; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                   (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beigelman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ID NO 944;
                                                                                                                                                                                         DNA; 15
                                                                                                                                                                                                                                                                                                                                                             0.6%;
                                                                                                                     entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                164pp; English.
                                                                                                                                                                                                                                                                            17
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Pred.
                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                               No. 57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Haeberli P,
                                                                                                                                                                                                                                                                                                                                                                               BB
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                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                            Gaps
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Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                      WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;
  Example 8; Page 85;
                                                                                                                                                                                                                                          Wraight
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                               21-JUN-2000; 2000WO-AU000693.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens.
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د
                                                                                                                                                                                                                                                                                         MURDOCH CHILDRENS RES
                                                                                                                                                                                                                                     Werther GA,
                                                                                                                                                                                                                                                                                                                                              99US-0140345P.
201pp; English
                                                                                                                                                                                                                                        Edmondson
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The present invention relates to a method for ameliorating the effects of CC skin disorders. The method comprises contacting the skin with an CC antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFB9]-2 or IGFB9], which is capable of inhibiting or reducing growth factor mediated cell proliferation, CC inhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense CC oligonucleotides of the present invention (see AAF45151 and AAF45153-CF F45161). The method is useful for ameliorating the effects of psoriasis, CC inthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC neoplasias, scleroderma, warts, benign growths, cancers of the skin, a CC hyperneovascular condition such as a neovascular condition of the retina, CC brain or skin, growth factor-mediated malignancies, other sclerotic CC disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia Sequence 15 BP; 2 A; 3 C; 6 G; 4 T; 0 Ф; 0 Other;

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WANTED THE TERM OF THE TERM OF
                                                                      Query Match
Best Local S
Matches 15
1232 GCATGTGCTGGCAGT 1246
                                                                      l Similarity
15; Conservat
                                                                      Conservative
                                                                                                   0.6%;
                                                                  0
                                                                                                       Score 15;
Pred. No.
                                                                      Mismatches
                                                                                                       36;
                                                                                                                                     DB 1;
                                                                  0
                                                                                                                                     Length 15;
                                                                      Indels
                                                                  0,
                                                                      Gaps
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GCATGTGCTGGCAGT 15

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RESULT 95
AAF02814/c
ID AAF028
XX AAF028
XX AAF028
XX I6-PBE
XX Hammer
XX KW Ribozy
KW interf
XX XX
XX XX
XX W02000
PD 19-OCT
                                                          Ribozyme; erythropoietin; interferon alpha; ss.
                                                                                                              16-FEB-2001
                                                                                                                                  AAF02814;
                                                                                                                                                     AAF02814 standard; DNA; 17
                                                                                         ribozyme substrate #1109.
                                                                                                             (first
                                                                                                             entry)
                                                                    granulocyte
                                                                    colony
                                                                    stimulating
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WO200061729-A2

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AAZ22430/

ID AAZ2

XX AC AAZ2

XX AC AAZ2

DT 25-N

XX Hum

XW Hum

XW Hum

XW bre

XX Syn

OS Syn

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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GAIA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythopoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                 Claim 3; Col 28; 24pp; English.
                                                                                                                         Antisense
                                                                                                                                                                                                                           Coswert
                                                                                                                                                                                                                                                                                                                                                                 18-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                  05-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5962672-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; RhoB protein; antisense oligonucleotide; disease; RhoB expression; breast cancer; primer; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ22430;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 0 A; 9 C; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 37; Page 81; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blatt L,
                                                                                                                                                                                                                                                                       (ISIS-)
                                                                                                                                                                                                                                                                                                                   18-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide directed against human RhoB mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ22430 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interferon alpha
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                                                                                                                                                                          1999-571296/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
15; Conserv
                                                                                                                                                                                                                                                                    SISI
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                                                                                               inhibition associated w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGAGGAGGACGAGGA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGAGGAGGACGAGGA 1897
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zwick M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                       PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
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                                                                                                  n of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.6%;
                                                                                               the gene encoding
n RhoB expression (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
51;
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                                                                                            g RhoB, usefo
e.g. breast
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                                                                                                                         useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                  cancer
                                                                                                                    for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>
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AAZ22392-Z22431 represent antisense oligonucleotides,

which

are

8-30

Query Match 0.6%; Best Local Similarity 88.9%; Matches 16; Conservative

0; Mismatches

<u>ب</u>

Indels

0,

Gaps

0

Score 14.8; Pred. No. 6

BB

<u>ب</u>

Length 18;

Sequence 18

BP; 4 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

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RESULT 97
RAF94683/c
ID AAF946
XX AAF946
XX AAF946
XX Rho; G
XX Rh
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                        This invention relates to an antisense compound targeted to a nucleic acid molecule encoding a member of the human Rho family of small GTP binding proteins, where the antisense compound inhibits the expression of the member of the human Rho family. The invention includes antisense coligonucleotides AAF94680 - AAF94637 which target a RhoA nucleotide sequence, AAF94725 - AAF94684 which target a RhoB nucleotide sequence, AAF94725 which target a RhoC nucleotide sequence, AAF94727 - AAF94766 which target RhoG nucleotide sequence, AAF94796 - AAF94790 which target a Rac 1 nucleotide sequence and AAF94795 - AAF94790 which target a Rac 1 nucleotide sequence and AAF94795 - AAF94790 which target a RhoC nucleotide sequence, AAF94795 which target cac42 nucleotide sequence. The antisense compound is useful for treating pretryproliferative conditions, especially cancer, abnormal wound healing or clotting conditions and ischaemia/reperfusion or recoxygenation injury. The compound may also be used to diagnose the above conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rho; GTP binding protein; phosphorothicate antisense oligonuclectide; RhoA; RhoB; RhoC; RhoG; Rac 1; cdc42; hyperproliferative condition; cancer; wound healing; clotting; ischaemia; reperfusion; reoxygenation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleotides in length, and are targeted to the gene encoding human The antisense oligonucleotides may be useful for treating diseases associated with the expression of RhoB, such as breast cancer. They also have research and diagnostic applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             An antisense compound targeted to a nucleic acid molecule encoding a member of the human Rho family of small GTP binding proteins useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roberts ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rho B antisense phosphorothioate oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-AUG-2000; 2000WO-US022808.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1251 CGGCTGCAGCAACAGCTG 1268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTP binding protein; phosphorothioate; RhoB; RhoC; RhoG; Rac 1; cdc42; hyper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001-191677/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 4 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 65; 156pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer and ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9908-00387341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
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RESULT 98
ABK50 27
ID ABK50
XX AC ABK50
XX Gepha
XW Cepha
XW Cepha
XW CahB;
XX WC200
XX WFI;
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                                                      RESULT 99
ADA27360
ID ADA27
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                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to production of Acremonium chrysogenum cephalosporin C-acetyl hydrolase (CPC-AH) and its utilisation in the synthesis of deacetylated derivatives of cephalosporin C (CPC) and the inactivation of the gene for increasing production of cephalosporin. Derivatives and/or their synthesis intermediates can be synthesised by growing a microbial host transformed with a DNA sequence that includes the cahB gene encoding A. chrysogenum CPC-AH, under conditions where it is either expressed or inactivated. The genes and proteins are used for removal of acetyl groups, especially from the 3'-carbon of CPC or from 7-aminocephalosporanic acid, to give deacetylated products useful as intermediates for cephalosporin antibiotics. Inactivation of the gene that expresses CPC increases production of cephalosporins by A. chrysogenum. This sequence represents a PCR primer used to clone the cahB gene encoding A. chrysogenum CPC-AH
ADA27360;
                                                      ADA27360 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 2 A; 4 C; 11 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Microbial production of cephalosporin C or its derivatives, useful as intermediates for antibiotics, in cells transformed with a gene encoding CPC (cephalosporin C)-acetylhydrolase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Velasco Alvarez J, Gutierrez Martin S, Casqueiro Blanco FJ;
Campoy Garcia S, Fierro Fierro F, Barredo Fuente JL, Diez Garcia
Martin Martin JF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cephalosporin C-acetyl hydrolase; CPC-AH; cephalosporin C; cahB; 7-aminocephalosporanic acid; ss; PCR.
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                                                                                                                                                                                                                                                                                                               Local 16;
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                                                                                                                                                                                                                                                                                                                               Similarity 88.5
16; Conservative
                                                                                                                                                                                                                                                                     GAGGCCGGCGAGGTGGAG 1838
                                                                                                                                                                                                              GAGGCCGGCGACGTGGGG 18
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                                                          DNA,
                                                                                                                                                                                                                                                                                                                                                            0.6%;
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Pred. No. 65;
0; Mismatches
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ADS78536
ID ADB785
AC ADB785
AC ADB785
XX O4-DEC
XX O4-DEC
XX Probe
XX Probe
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XX Synthe
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                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an oligonucleotide primer capable of specifically hybridising to a DNA having the sequence of the flanking regions of a microsatellite selected from M2-4-9, M2-2-12, M2-3-11, M2-2-20, M2-2-21, M2-2-22, M2-2-23, M2-2-24, M2-4-25, M2-4-26, M2-2-29, M2-2-32, M2-4-32, M2-4-33, M2-4-37, M2-3-22, M2-2-36, M2-5-11, M2-2-46, and M2-2-48. The primer is useful for determining the number of repeat units of the microsatellite cited above. The primer is useful in HLA-related research, such as genetic mapping of HLA class II-associated diseases, transplantation matching, population genetics, and identification of recombination hot spots as well as linkage disequilibrium studies. The present sequence represents the human microsatellite repeat M2_3_4.
                                                                                                        human leukocyte
                                                                          Synthetic
                                                                                                                                    Probe sequence #39 related to the invention.
                                                                                                                                                                04-DEC-2003
                                                                                                                                                                                              ADB78536;
                                                                                                                                                                                                                           ADB78536 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 7 A; 0 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotide primer capable of specifically hybridizing to a DNA having the sequence of the flanking regions of a microsatellite (e.g. M249), useful for HLA-related research, e.g. transplantation matching.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ds; HLA-related research; HLA class II-associated disease;
transplantation matching; recombination hot spot identification;
linkage disequilibrium study; human; microsatellite.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human microsatellite repeat M2_3_4.
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                                                                                                                                                                                                                                                                                                                              1770 GAGGAGGAGGAGGCGGAG 1787
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16; Conservat
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                                                                                                                                                                  (first
                                                                                                        antigen;
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                                                                                                                                                               entry)
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                                                                                                        HLA;
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Pred. No. 6
                                                                                                        probe;
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03-APR-2003 WO2003027309-A2

24-SEP-2001; 2001US-0324421P

24-SEP-2002;

2002WO-US030238

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RESULT 101
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Best Local S
Matches 16
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          WPI; 2002-089915/12
                                                                                                                        23-MAY-2001; 2001WO-US016908
                                                                                                                                                                               WO200190124-A2
                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                   Human; G-protein coupled receptor 31; GPR31 protein; haplotyping; genotyping; gene therapy; cancer; polymorphism; ASO; primer; allele-specific oligonucleotide; ss.
                                                                                                                                                                                                                                                                                            Human GPR31
                                                                                                                                                                                                                                                                                                                     26-MAR-2002
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                                                                                          23-MAY-2000; 2000US-0206572P
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                                                                  (GENA-) GENAISSANCE PHARM INC
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                                                                                                                                                                                                                                                                                                                     (first entry)
                                      Duda
                                                                                                                                                                                                                                                                                         polymorphism detecting ASO primer #16
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                                        Kazemi
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Pred. No. 65;
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                                        Lee
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                                        Messer C;
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Novel genetic variants of G-protein coupled receptor studying expression and function of the protein, and to treat diseases e.g. cancer. gene useful in for screening drugs

Claim 16; Page 13; 75pp; English

receptor 31 (GPR31) gene. The invention also relates to compositions and methods for haplotyping and/or genotyping the GPR31 gene in an individual. Polynuclectides of the invention are useful in studying the expression and function of GPR31, and in expressing GPR31 protein for use in screening candidate drugs to treat diseases related to GPR31 activity of and in studying the effect of the variation on the biological activity of GPR31 as well as on the binding affinity of candidate drugs targetting GPR31 as well as on the binding affinity of candidate drugs targetting GPR31 for the treatment of cancer. They are also used in gene therapy. The haplotyping method is useful for improving the efficiency and creliability of several steps in the discovery and development of drugs for treating diseases associated with GPR31 gene in an individual, which can also be used by the pharmaceutical research scientist to validate GPR31 as a candidate target for, and in design of clinical trials of candidate drugs, for treating a specific condition drugs or disease predicted to be associated with GPR31 activity. The present sequence is an allele specific oligonucleotide (ASO) primer used to detect human GPR31 gene polymorphisms The invention relates to genetic variants of human G-protein coupled GPR31) gene. The invention also relates to compositions for use e E

Sequence 15 BP; 4 A; 3 C; 6 G; 1 T; 0 U; 1 Other;

밁 Ś Query Match Best Local S Matches 14 2106 ¢G¢ŢŢ¢¢ŢĠ¢Ā¢ 2120 15 14; Similarity CRCTTCCTGCTGGAC 1 Conservative 93.3%; 0.6%; <u>,</u> Score 14.6; Pred. No. 44 Mismatches DB 1; Length 15; 0 0 Gaps 0

RESULT 102

AAT76198 standard; DNA; 16 ВP

AAT76198

12-SEP-1997 (first entry)

Human IL4 receptor antisense oligonucleotide

Asthma; airway epithelium; adenosine free; cystic fibrosis; chronic obstructive pulmonary disease; bronchitis; interleukin; ss

Synthetic

WO9640162-A1

19-DEC-1996

06-JUN-1996; 96WO-US009306

07-JUN-1995; 95US-00474497

(UYEC-) UNIV EAST CAROLINA

Ä Metzger WJ;

WPI; 1997-051871/05

AAT76198/c
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A Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of

Example 5; Page 30; 71pp; English

A method for treating airway disease in a subject has been produced

which involves the topical administration of an essentially adenosine

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RESULT 103
ABK67989
ID ABK679
XX ABK679
XX ABK679
XX Mutant
XX Mutant
XX Mutant
XX Mutant
XX Mutant
XX Mo200;
XX Synth
XX W0200;
XX W0200;
XX W0200;
XX W0200;
XX W71
PD 04-API
XX (MITU
PF 26-SEI
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PF 26-FEI
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PF 27-FEI
PF 20-FEI
PF 20-FEI
CC DNA 1
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Best Local S
Matches 15
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                        The present invention relates to a new method of constructing a mutant DNA library. The method of the invention involves ligating unit domain DNAs in arbitrary combinations, mixing the ligated unit domain and performing polymerase chain reaction (PCR) by employing the ligated unit domain DNA mixture as a template to obtain a DNA library containing 2 or more clones. The method can be used for constructing a mutant DNA library which is for use in producing e.g. protein library, mutant proteins and artificial amino acid sequences with desirable functional properties with a combination of unit domains. The method can also be used for selectively splicing a library to yield mutant proteins with retention of function and smaller for wider application and in evolution molecular engineering in which the constructed mutant DNA library contains various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Constructing mutant DNA library comprises ligating unit domain DNAs in arbitrary combinations before mixing in specific manner as template for polymerase chain reaction to give clones, useful in producing protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mutant DNA library; selective splicing;
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06-FEB-2001; 2001JP-00029138.
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   including
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unit domain; clone; protein library; mutant protein; molecular engineering; DNA shuffling; evolution; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry.
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       original
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Pred. No. 57;
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              exons and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 16;
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Query Match Best Local S Matches 10

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Score 14.4; D Pred. No. 67; 5; Mismatches

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RESULT 104
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Best Local
                     The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fins-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           regulating domains in structural DNAs. The method is easy and less error prone in DNA shuffling when constructing DNAs. The present nucleic acid sequence represent one of a collection (ABK67951-ABK68000) of PCR primer; that were used in the methods of the invention for construction of a
                                                                                                                                                                                                                Nucleic acid molecule modulating VEGF receptor(s) gene expensibility - useful for treating e.g. tumour angiogenesis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                              26-OCT-1995;
11-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular fms-like tyrosine kinase 1; kinase insert domain containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX72692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mutant DNA library, as describe above
                                                                                                                                                                                       Claim 4; Page 126; 218pp; English.
                                                                                                                                                                                                                                                                            HPI;
                                                                                                                                                                                                                                                                                                        Pavco
                                                                                                                                                                                                                                                                                                                                    (RIBO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; RNA; 17
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                                                                                                                                                                                                                                                                                                        Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VEGF receptor hammerhead ribozyme substrate
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96US-00584040.
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Pred. No. 57;
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                                                                                                                                                                                                                                                 gene expression
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RESULT 105
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A method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endounclease activity and catalytic activity; from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic ascides and infection. They may also be used to detect genetic drift and mutations in diseased cells and to determine c-raf RNA. Specifically NACs with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restensels, soriasis or rheumatoid arthritis, or generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity. ANY9922 to ANY9387 represent NACs that can be used in the method, specifically for modulating the expression of a Raf gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying new catalytic nucleic acid especially ribozymes that cleave Raf restenosis, and also new ribozymes and used as antiviral agents and synthons.
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03-JUL-1997;
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02-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-OCT-1997;
05-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  target; substrate; catalyst; modulation; expression;
screening; identification; synthesis; deprotection; 
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Beigelman L, Mcswiggen
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97US-0051718P.
97US-0056808P.
97US-0061321P.
97US-0061324P.
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Sweedler D;
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids may be enzymatic nucleic acids (e.g. a riboxyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) presence of a divalent cation that is preferably Mg^2+.
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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                                                                                                                                                                                                                                                                                   88; Page 131; 200pp; English.
                                                                                                                                                                                                                                                                                                                      L actu motecules, e.g., enzymatic nucleic acids and antisense ucts, which down regulate expression of a CD20 gene or neurite inhibitor gene useful for treating, e.g., lymphoma, leukemia, l nervous system injury.
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06-MAR-2000;
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    (RIBO-)
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    RIBOZYME PHARM INC.
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2000US-0187128P
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Claim 88; Page 131; 200pp; English
                                                                                                                                                                                                    Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, central nervous system injury.
                                                                                                                                                                                                                                                       (CHOW/) CHOWRIRA B
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The invention relates to a nucleic acid molecule which down regulates CC expression of a CD20 gene and a nucleic acid molecule which down considered to the contact of a neurite growth inhibitor gene (NGGO). The CC mucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CD NAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an RNA motif), a G-cleaver (cleaving RNA with a NGN motif) processing an RNA with a NGN triplet), a zinzyme (cleaving RNA with a CD20 targetting RNA with a cell to reduce CD20 activity of the CD20 in the presence of a divalent having a condition associated with the level of CD20. The treatment may further comprise the use of one or more condition associated with the level of lymphoma (NGL), immunocytoma (NGL), small B-cell lymphocytic lymphoma, (NGL), immune thrombocytopaenia, and inflammatory arthropathy. The NGCO-targetting nucleic acid may be contacted with a cell to reduce NGGO gene in the contacted acid may be contacted with a cell to reduce NGGO activity of the NGCO-therapies. In particular, the NGGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke), Alzheimer's disease, dementia, multiple sclerosis (NS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (NLS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (NLS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (NLS), chemotherapy-induced neuropathy.

CC interest controlled to the

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17
BÞ;
7
A,
0
C; 10 G; 0 T; 0 U;
0 Other;
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Query Match Best Local 15; Similarity GGAGGAGGACGAGGAG 1898 Conservative 0.6%; 93.8%; 0 Score 14.4; Pred. No. 67; Mismatches DB <u>,</u> 1; Length 17; Indels 0 0

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           S
                         Matches
N
GGAGGAGGAAGAGGAG 17
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RESULT 108
ABA78078/C
ID ABA780
XX ABA780
XX ABA780
XX ABA780
XX ABA780
XX ABA780
XX HUMAN;
XX HUMAN;
XW YETINC
XW YETINC
XW CYCILT
XW Adenom
XW Adenom
XW Mismat
XW Mismat
XW Mismat
XW Famili ABA78078, ABA78078 standard; DNA; 17 (first entry) ВP

mismatch repair; MSH2; MSH6; hy familial hypercholesterolaemia; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoplobin alpha tocus 1; MEHI; AFOE mismatch repair; MSH3; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGTI; syndrome; APP; PSENI; antisense; Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; Cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; Cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; Cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; Cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; Cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; CDKN2A; melanoma; APC; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; CYClin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; CYClin-dependent kinase inhibitor 2A; CYClin-dependent ki correcting oligonucleotide SEQ ID NO: 924. APOE;

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RESULT 109
ABA78077
ID ABA780
XX ABA780
AC ABA780
XX BACA
DT 24-JAN
XX BE BRCA1
XX PSE BRCA1
XX PSE STAIN
XX M CYClin
XW retinc
XW Ademor
XW Ademor
XW Ademor
XW Ademor
XW Ademor
XW Ademor
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                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 15
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27-MAR-2000;
01-JUN-2000;
30-OCT-2000;
Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CPTR; cystlo fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                               BRCA1 mutation
                                                                                                                                                24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
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                                                                                                                                                                                                              standard; DNA; 17
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; 2000US-0192179P.
; 2000US-0208538P.
; 2000US-0244989P.
                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 1
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                                                                                                                                                (first
                                                                                                               correcting oligonucleotide SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                       A.
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                                                                                                                                                                                                                                                                                                                                                                                                                       6 C; 3
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0: Mismatches
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Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                         U; 0 Other;
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                                                                                                                 923
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RESULT 110
ABA78070/c
ID ABA780
AC ABA780
AC ABA780
XX 24-JAN
XX 24-JAN
XX ETIL
CKW FUMAN;
KW retil
KW

ABA78070 ABA78070

standard;

DNA;

BRCA1 mutation

correcting

oligonucleotide SEQ

ID NO:

916.

entry)

Human; gene therapy; adenosine deaminase deficiency; positional rethroblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; APCB;

S

GCAGCAACAGCTGGAA 1271 GGAGCAACAGCTGGAA

Query Match Best Local S Matches 15

Similarity

93.8%;

Conservative

0

Score 14.4; Di Pred. No. 67; 0; Mismatches

B

1; Length 17;

0

Gaps

0

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The present invention provides single-stranded oligonucleotides which can compute the used for the targeted alteration of genomic sequences, where the coligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MIH, MSH2, MSH6, apolipoprotein E (APOB), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APC), presenilin-1 (PSEN1) and cresenilin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, such as cancer, adenosine deaminase deficiency, cystic fibrosis, alpha locus syndromes. The present sequence is one of the gene correcting coligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide for targeted alterations of genetic sequences treating cystic fibrosis, comprises at least one mismatch and modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo
Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAR-2000; 2000US-0192176P.
27-MAR-2000; 2000US-01921799.
01-JUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            familial hypercholesterolaemia; UGT1; syndrome; APP; UDP-glucuronosyltransferase; amyloid precursor protei Alzheimer's disease; cytostatic; antisickling; antiar
                                                                                                                                                                                                                                                                                                                                           Claim 7; Page 100;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antilipemic; ss.
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BP; 7 A; 3 C; 6 G; 1
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N
T;
U; 0 Other
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                                                                                                                                                                                                                                                                                                                                                                                              and for
chemical
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RESULT 111
ABA78069
ID ABA7806
XX ABA780
XZ ABA780
XX BBA780
XX BBA780
XX ETAIN
XX FURING
XX BRCA1
XX FURING
XW FURING
KW Cyclin
KW Cyclin
KW Gyclin
KW Gyclin
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                                                                                                                                                                                                                                                                                                      5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the coligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p33, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Factor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Factor VIII, Factor III, MSH2, MSH6, apolipoprotein S (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APC), presenilin-1 (PSEM1) and presenilin-2 (PSEM2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting coligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 15
retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
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27-MAR-2000; 2000US-0192179P.
01-JUN-2000; 2000US-0206538P.
30-OCT-2000; 2000US-0244989P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEM1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide treating cystic
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                                                                                     BRCA1 mutation
                                                                                                                                                                                             ABA78069 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 1 A; 6 C; 3 G; 7 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modification
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                                                                                                                                                                                                                                                                                                                                                   15;
                                                  gene therapy;
                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                 correcting
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                                     deaminase deficiency; p53; beta-globin;
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RESULT 112
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XX Human
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29-MAY-2002 ABN00936;

(first 17-mer

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sequence

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hGDMLP-1; heart;

ABN00936 standard;

DNA;

17

Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.

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GGAGCAACAGCTGGAA 17

1256 ССАССААСАССТССВА 1271

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Query Match
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Matches 15
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27-MAR-2000;
01-JUN-2000;
30-OCT-2000;
                                                                                            oligonuclectide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKNJA), APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein B (APCB), LDL receptor (LDLB), UDP-glucuronosyltransferase (UGT1), amyloid precursor protein (APC), presenilin-1 (PSEN1) and presenilin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemoghilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous spolyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                   Claim 7; Page 99; 294pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide for targeted treating cystic fibrosis, com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
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l Similarity
15; Conserv
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Conservative
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d. No. 67;
Mismatches
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                                                                                                                                                                   cc expressing the proteins. The hGDMLP-1 proteins or polypeptides may be expressing the proteins. The hGDMLP-1 proteins or polypeptides may be cused as immunogens to raise antibodies that specifically recognise hGDMLP countries. As standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule coupture probes for surface-enhanced laser description ionisation, as the representation and in vaccines or for replacement therapy in hGDMLP-1 production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The conjunct of the sequences encoding hGDMLP-1 may be used for diagnosing a conjunct of the sequence sencoding hGDMLP-1 is localised to chromosome 22.

Cr and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

Cr he present sequence represents an oligomer used in the screening of the present invention. N.B.

Cr he sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO can be sequenced as a companion of the printed specification, but was obtained in electronic format directly from WIPO can be supposed to the printed specification.
                                                         Query Match
Best Local S
Matches 15
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21-SEP-2000;
27-SEP-2000;
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                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 928; 214pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser protein ionization, comprises human myosin-like protein hGDMLP-1.
                              1264 AGCTGGAAGAGGCTGA
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RESULT 113 ABN02626

Sequence 17 BP; 5 A; 4 C; 6 G;

2 T; 0 U; 0 Other;

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cc nucleic acids can be used as probes to detect, characterise and quantify cc hgbMLp-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hcbMLp-1 cc protein variants having desired phenotypic improvements, and for ce expressing the proteins. The hgbMLp-1 proteins or polypeptides may be comed as immunogens to raise antibodies that specifically recognise hcbMLP-1 proteins, as standards in assays used to determine the concentration cc and/or amount specifically of hcbMLp proteins, as specific biomolacule capture probes for surface-enhanced laser desorption ionisation, as the repeated of the protein and in vaccines or for replacement therapy. The concentration cc production, and in vaccines or for replacement therapy. The concentration alsorder associated with the expression of hcbMLP-1, in particular heart cand skeletal muscle disorders. hcbMLp-1 may be used for diagnosing a client sequence represents an oligomer used in the screening of the hcbMLP-1 sequence in the exemplification of the present invention. N.B. cr hcbMLp-1 sequence attacked in electronic format directly from WIFO at ftp. wipo.int/pub/published_pct_sequence
26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                                                                                                                                                             he present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-can be used in gene therapy and vaccine production. The hGDMLP-1
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The
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27-SEP-2000;
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as specific biomolecule capture probes for surface-enhanced laser
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93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                         8658; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprises human myosin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hanzel
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Pred. No. 67;
0; Mismatches
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RESULT 115
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Human;
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Best Local &
Matches 15
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30-JAN-2001;
30-JAN-2001;
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21-SEP-2000;
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30-JAN-2001;
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2001WO-US000661.
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2000US-0234687P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chromosome 22; gene therapy; vaccine; heart disease; disorder; amplicon; screening; ss.
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93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO:5 sequence SEQ ID NO:6392.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of 1 can be used in gene therapy and vaccine production. The hGDMLP-1

hGDMLP-

Disclosure; desorption

SEQ ID NO

6392;

214pp;

English

ionization,

e, for raising antibodies that recognize hGDMLP-1 proteins, biomolecule capture probes for surface-enhanced laser ization, comprises human myosin-like protein hGDMLP-1.

2002-179446/23.

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RESULT 116
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Matches 15
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30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
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21-SEP-2000; 2000US-0234687P.
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Pred. No. 6
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of a
                                                                                                                                              Disclosure; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-179446/23.
                                                                                                                                         6391; 214pp; English.
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CC protein 1 (hGDMLP-1). The protein and polymuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 cc nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as capture probes for surface-enhanced laser description ionisation, as colution, and in vaccines or for replacement therapy. The proteins and in vaccines or for replacement therapy. The copynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a cliented with the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present invention. N.B. Chester of the protein of the present invention. N.B. Chester of the protein of the present invention. N.B. Chester of the protein of the protein of the protein of the protein of the printed capture. The sequence data for this pattent did not form part of the printed capture in the samplification of the present invention. N.B. Chester of the printed capture of the printed capture of the printed capture. The sequence data for this pattent did not form part of the printed capture of the printed capture. The sequence data for this pattent did not form part of the printed capture. The sequence data for this pattent did not form part of the printed capture. BP; 3 A; 8 C; 4 G; 2 T; 0 U; 0 Other;

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              8
                                  Matches
                                         Query Match
Best Local (
                                                                Sequence 17
                                        Local
               1602 GCCCCGTGCTTCCAGA 1617
N
                                 l Similarity
GCCCCGTGCTCCCAGA 17
                                  Conservative
                                         93.8%;
                                 0;
                                        Score 14.4;
Pred. No. 67;
                                 Mismatches
                                                 DB 1; Length 17;
                                   Indels
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                                   Gaps
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0

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RESULT 117
ABN02625
ID ABN026
XX ABN026
XX ABN026
XX ABN026
XX Human
DE Human
XX Human
XX Human
XX Human
XX Homo s
XX W02001
XX W02001
XX W02001
XX W201
XX W2001
XX W2001
XX W2001
XX W2001
XX W02001
XX W020
26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
04-CCT-2000;
30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
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    ; 2000US-0207456P.

2000US-0234567P.

2000US-0236359P.

2000GB-00024263.

2001WO-US000661.

2001WO-US000663.

2001WO-US000663.

2001WO-US000663.

2001WO-US000665.
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heart disease;
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RESULT 118
ABNO09
XX ABNO09
XX ABNO09
XX ABNO09
XX ABNO09
XX Human;
XX Human;
XX Human;
XW muscle
XW skelet
XX Homo s
XX Homo 
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Best Local
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30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
                                                                                                                                                                                                                                                  Human, genome-derived myosin-like protein 1, GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
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                                                                                                                                                                                                                                                                                                                                                                                                    scanning
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                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO:4 sequence SEQ
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                                                                                                                                                                                                                                                                                                hGDMLP-1; heart;
heart disease;
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RESULT 119
ABNO8668
ID ABNO86
XX
AC ABNO86
XX
DT 29-MAY
XX
DE Human

standard; DNA; 17

0

29-MAY-2002 ABN08668; ABN08668

(first entry)

Human GDMLP-1 17-mer scanning

SEQ

IJ.

NO:5

sequence

SEQ

H

NO:8660

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                                                                                                                                                                              The present invention describes a human genome-derived myosin-like CC protein 1 (hdDMLP-1). The protein and polynucleotide sequences of hdDMLP-1 CC I can be used in gene therapy and vaccine production. The hdDMLP-1 CC mucleic acids can be used as probes to detect, characterise and quantify CC hdDMLP-1 nucleic acids in sample, as amplification substrates, to provide initial substrates for the recombinant engineering of hdDMLP-1 CC protein variants having desired phenotypic improvements, and for cexpressing the proteins. The hdDMLP-1 proteins or polypeptides may be comed as immunogens to raise antibodies that specifically recognise hdDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hdDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as the capture probes for surface-enhanced laser description ionisation, as complement in pattients having specific deficiency in hdDMLP-1 capture probes for surface-enhanced laser description ionisation, as complement in pattients having specific deficiency in hdDMLP-1 composited with the expression of hdDMLP-1 in particular heart conditions and skeletal muscle disorders. hdDMLP-1 may be used for diagnosing a complement sequence represents an oligomer used in the screening of the present sequence in the expression of hdDMLP-1 in particular heart complement in patticular heart complement in the sequence in the screening of the present sequence data for this patent did not form part of the printed capture when into the millished not fermence.
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Matches 15
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21-SEP-2000;
27-SEP-2000;
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30-JAN-2001;
30-JAN-2001;
                                                                                                                                  Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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30-JAN-2001;
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                                 1265 GCTGGAAGAGGCTGAG 1280
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2000US-0234585P.
2000US-0236359P.
2001WO-US000661.
2001WO-US000662.
2001WO-US000664.
2001WO-US000665.
2001WO-US000665.
2001WO-US000667.
2001WO-US000667.
                                                                                                                                    BP; 5 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
                                                                  Conservative
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                                                                  Score 14.4; D
Pred. No. 67;
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hGDMLP-1; heart;

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                                                                                                                                                                               therapeutic supplement in patients having specific deficiency in hgDMLP-i production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hgDMLP-1 may be used for diagnosing a disorder associated with the expression of hgDMLP-1, in particular heart and skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hgDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; hear muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                   Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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30-JAN-2001;
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27-SEF-2000;
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30-JAN-2001;
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                                                                                                                                                                  ftp.wipo.int/pub/published_pct_sequence
                                1530
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                                                                                  Similarity
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                                                                                                                                   BP; 7
                                                                 Conservative
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2001WO-US000667.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INC.
                                                                                                                                   A; 3 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                0.6%;
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                                                               Score 14.4; I
Pred. No. 67;
0; Mismatches
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                                                                                                DB 1;
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                                                                                                Length 17;
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EP1229046-A2

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RESULT 120
AAD45173/c
ID AAD451
XX AAD451
AC AAD451
AC AAD451
XX 27-DEC
XX 27-DEC
XX Primer
XX Human;
KW Primer
XX US6426
XX US6426
XX 01-AUC
XX 0
ABV7892
ID ABV7892
ID AB
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AC AB
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XX HU
XX PI
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Best Local S
Matches 15
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AAD45173
                                                                                                                                                                                                      Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; ktdney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to antisense compounds targetted to a nucleic acid encoding human receptor interacting protein (RIP)2 to inhibit its expression. Antisense compounds are used for treating diseases associated with RIP2 expression. They are also useful in antisense gene therapy. The present sequence is human RIP2 DNA specific PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 4 A; 4 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotide that targets regions of a encoding human receptor interacting protein (RIP)2, for associated with RIP2 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABV78926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-AUG-2001; 2001US-00920663.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human RIP2 DNA specific forward PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABV78926;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Human HTPL scanning
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCTTCACGGCCACCTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.4;
Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   67;
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treating diseases
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ABV7/
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                                                                                                                                                                                                                                                                   RESULT 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to human testis expressed Patched like protein (HTPL, see ABV875) to ABV78762 and ABB98519 to ABB98520). HTPL CC has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL CC shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is CC important in regulating male germ cell development, and the HTPL gene was mapped to human chromesome logle 1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in CC therapy and manufacture of a medicament for treatment or prevention of CC such disorder associated with decreased expression or activity of human CC HTPL. Such disorders include disorders of testis, or adrenal, adult and CC foetal liver, bone marrow, brain, kidney, lung, placenta, prosstate, collanding the collon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an examinate from the invention.
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                          Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 2 A; 6 C; 7
                                                                                                                                Human HTPL scanning
                                                                                                                                                                                                  ABV78924;
                                                                                                                                                                                                                                 ABV78924 standard; DNA; 17
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2001WO-US000664.
2001WO-US000665.
2001WO-US000667.
2001WO-US000668.
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2001US-00864761.
2001US-0327898P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the invention
                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                     0.6%;
93.8%;
                                                                                                                                                                entry)
                                                                                                                                oligonucleotide SEQ ID 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.4;
Pred. No. 67;
                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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RESULT 123
ABK17396/C
ID ABK173
XX ABK173
XX O9-AFK173
XX O9-AFR
XX W O9-AFR
XX W Ophtha
KW Vulner
KW Vulner
KW Uner

ERG

hammerhead ribozyme target sequence,

Seq

ij

ö 43 09-APR-2002

(first entry)

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopatl vulnerary; cancer; lymphoma; Ewing's Barcoma; melanoma; psoriasis; tumour anglogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgi

verruca vulgaris;

osteopathic;

7396/C ABK17396

standard; RNA;

17

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                        S
                                                                                                                                                            The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL CC has two isoforms, with a few single base pair differences between the CC two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL CC shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is CC important in regulating male germ cell development, and the HTPL gene was CC import to human chromosome 10pl2.1. HTPL and its coding sequence are CC useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of CC therapy and manufacture of a medicament for treatment or prevention of CC such disorder associated with decreased expression or activity of human CC foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, CC skeletal muscle or colon function. HTPL proteins and nucleic acids are CC clinically useful diagnostic markers and potenial therapeutic agents for CC manned from the invention.
                                                             Best Loc
Matches
                                                                                         Query Match
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23-MAY-2001;
09-CCT-2001;
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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-676582/73.
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                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AEOM-)
                                                                          Local
                             2120 CCACGGGGGCGGAGTG 2135
                                                             l Similarity
15; Conser
N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AEOMICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Page 86; 718pp; English.
                                                                                                                                                       from the invention
 CCACCGGGCCGCAGTG
                                                         0.6%;
ilarity 93.8%;
Conservative (
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2001WO-US000664.
2001WO-US000667.
2001WO-US000667.
2001WO-US000668.
2001WO-US000669.
2001US-00864761.
2001US-00327898P.
                                                                                                                      BP; 2 A; 7 C; 7
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   17
                                                                                                                        G; 1
                                                             0,
                                                                                          Score 14.4;
                                                                             Pred.
                                                           red. No. 67;
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                                                                                            DB 1;
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                                                                                          Length 17;
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RESULT 124
ABK17919/c
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                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 15
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Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss;
Osler-Weber-rendu syndrome, leukaemia; ostsoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
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                 09-APR-2002
                                                       ABK17919;
                                                                                                ABK17919
                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RIBOZYME PHARM INC. GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                            BP; 1
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               (first entry)
                                                                                                                                                                                                                                                                                                                                                          A; 9
                                                                                              RNA; 17
                                                                                                                                                                                                                                                                                                0.6%;
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                                                                                                                                                                                                                                                                                                Score 14.4;
Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                            U; 0 Other;
                                                                                                                                                                                                                                                                                                       67
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vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulga angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; osteoporosis; DNAzyme; inozy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; hammerhead ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ERG hammerhead ribozyme target sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hammerhead
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ribozyme; cytostatic; antitumour; antidiabetic; antiarthritic; antipsoriatic; virucide; osteopa
                                                                                                                                                                                                                                                                                                                                                                                                                   osteoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq
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WO200188124-A2 sapiens.

22-NOV-2001

16-MAY-2001; 2001WO-US015866

16-MAY-2000; 2000US-00572021.

(RIBO-) RIBOZYME PHARM INC. GLAXO GROUP LTD.

WPI; 2002-082995/11 Η, Mcswiggen JA, Mclaughlin F, Randi AM

Novel polynuclectide which down regulates expression of Ets-related gene useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

Claim 4; Page 69; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates CC expression of an Ets-related gene (ERG). (I) is useful for treating CC conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, CC tumour angiogenesis, diabetic retinopathy, macular degeneration, verruca CC neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca CC vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge CC weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu CC syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, CC the treating sells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies CC under conditions suitable for the treatment. Leukaemia or tumour CC angiogenesis is treated by administering (I) to the patient in CC early by contacting the cell with (I). (I) is useful for reducing ERG activity in a CC early by contacting the cell with (I). (I) is useful for leukaemia or tumour CC early such as Mg2+. (I) is useful for reducing ERG activity in a ccell by contacting the cell with (I). (I) is useful for cleaving RNA of CC examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically CC examine genetic drift and mutations within RRG gene or ERG fusion genes. CRS targeting genes that share homology with ERG gene or ERG fusion genes. CRS targeting antienesse and CC englated PCR primers of the invention

Sequence 17 BP; 2 A; 9 C; 3 φ; 0 .. H w 뎍

皮 Ś Query Match Best Local S Matches 15 17 15; Conservative CTGGAGGAGGACGCGG 2 CTGGAGGAGGACGAGG 1896 93.8%; 0; Score 14.4; D Pred. No. 67; 0; Mismatches DB 1. 0 Gaps

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ABT377829/c
ID ABT378
XX
AC ABT378
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AC ABT378
XX
AC ABT378
XX
Cytost
KW Cytost
KW Schizz
XX
Homo |
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WO2000
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PD 17-SE
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PD 17-SE
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PD 17-SE
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PD 77-SE
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INGLE
IN
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                                                                                                                                                                                                                                                                                                                                     Consider the specification, a sequence containing at least 15 consecutive controlles from the 17 mer sequence with, after optimal considered that least 80 % identity to the 17 mer sequence, a sequence that conditions, or the complement confidered to them under highly stringent conditions, or the complement conflictions, or the complement conflictions of the invention are useful as probes and primers for detecting, confidentifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for component of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell containing the sparastion, specifically cancer but also Alzheimer's disease and confidence to smalles is useful for diagnosis and/or prognosis of these confidence is a seeful for diagnosis and/or prognosis of these confidence is a seeful for diagnosis and/or prognosis of these confidence is a seeful for confidence and antibodies are useful as components of protein confidence. The mucleic acid sequences of the invention can be used in gene confidence in the polypeptide and antibodies are useful as components of protein confidence in the polypeptide and antibodies are useful as components of protein confidence in the polypeptide and antibodies are useful as components of protein confidence in the polypeptide and antibodies are useful as components of protein confidence in the polypeptide and antibodies are useful as components of protein confidence in the polypeptide and antibodies are useful as components of protein confidence in the polypeptide confidence represents a tumour suppression confidence in the polypeptide confidence of the invention can 
                                                                                                                                        Query Match
Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Telerman A, Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-2001; 2001FR-00011978
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                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MOLE-) MOLECULAR ENGINES LAB
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   16
                                                                                                                                        l Similarity
15; Conserv
                                             CTGCGTTTTGCTGAGC 1398
                                                                                                                                                                                                                                                                                        17
   CIGCGITITGCIGATC 1
                                                                                                                                                                                                                                                                             BP; 7
                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                A; 4 C; 5 G; 1 T; 0 U; 0 Other;
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                                                                                                                                    0;
                                                                                                                                                                              Score 14.4;
Pred. No. 67;
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                                                                                                                                            Mismatches
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ation; cancer; Alzheimer
                                                                                                                                                                                                        DB 1;
                                                                                                                                        1;
                                                                                                                                                                                                            Length
                                                                                                                                            Indels
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s disease;
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ55389 - ABZ65216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human riboxymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                 Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                           WPI;
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06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P
Sequence 17
                                                                                                                                                                                                                                                                   Claim 4; Page 139; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002; 2002WO-US016840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human HER2 DNAzyme substrate
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                                  ribozymes of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                           2003-140484/13
BP; 3 A; 7 C; 3 G; 0 T; 4 U; 0 Other;
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                                   invention
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                 Matches
                       Query Match
Best Local
   1266
                        Similarity
   CTGGAAGAGGCTGAGG 1281
                 Conservative
                       93.8%;
                 0
                       Score 14.4;
Pred. No. 67;
                 Mismatches
                               DB
                              1;
                 1;
                              Length 17;
                 Indels
                 0;
                 Gaps
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0

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17

CTGGAAGACGCTGAGG

N

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RESULT 127
AAA10553/c
ID AAA10553/c
XX AAA105
XX AAA105
XX PT 29-JUN
XX Smad2
XX Smad2
XX Smad2
XX Smad2
XX Chromc
XX diagnc
                                                                                                        AAA10553;
                                                                                                                             AAA10553
                                                         Smad2 antisense oligonucleotide sequence
                                                                                 29-JUN-2000
                                                                                                                               standard; DNA; 18
                                                                                 (first entry)
                                                          #6 (ISIS# 27783).
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Smad2; MADH2; MADR2; hMAD2; JV18-1; transcription factor; inflammation; chromosome 18q21; antisense compound; treat; prevent; infection; tumour diagnostic reagent; research reagent; ss; cancer.

RESULT 126 ABZ64881/c ID ABZ64881 standard;

RNA;

17

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RESULT 128
ABS60576/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents an antisense nucleotide sequence targeting human smad2. Smad2 is also known as MADH2, MADR2, hMAD2 and JVL8-1, and is a member of a subgroup of Smad family transcription factors which are cytosolic proteins regulated by transforming growth factor-beta (TGF-beta) and activins. Smads exist as monomers in unstimulated cells as home or heterodimerise and translocate to the nucleus and activate target gene transcription upon ligand binding. The Smad2 gene is located on chromosome 18q21. The invention relates to antisense compounds (see AAA10548-A10587) targeted to the Smad2 nucleotide sequence. The antisense aligonucleotide sequences inhibit Smad2 surression by hybridising to DNA or RNA. The antisense nucleotides are used to treat or prevent diseases associated with expression of Smad2, e.g. infection, inflammation and tumours. The oligonucleotides can also be used as diagnostic or research
                                                       Aminopeptidase P, XPNED2; bradykinin receptor B1; ds; BDXRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein KLKI; bradykinin receptor B2; BDXRB2; gene therapy; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; P14; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD; chronic obstructive pulmonary disease; enterocolitis.
                                                                                                                                                                                                                                                                                                                                                                          ABS60576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 11; Col 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound that inhibits human Smad2, useful e.g. treating or preventing infection, inflammation and tumors.
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                                                                                                                                                                                                                                                                         Human polymorphism associated DNA sequence #325.
                                                                                                                                                                                                                                                                                                            05-NOV-2002
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Pred. No. 78;
O; Mismatches
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WO200261131-A2

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AGAGCGGAGAGCGAAG ACAGCGGAGAGCGAAG 1815

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Query Match Best Local S Matches 15

l Similarity 93. 15; Conservative

93.8%;

Score 14.4; Pred. No. 78; Mismatches

DB 1; Length 18;

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Gaps

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CC encoding aminopeptidase P (KPNEP2), bradykinin receptor B1 (BDKRB1), C1 esterase inhibitor (C1NH), kalikrein CC (achykinin receptor B2 (BDKRB2), angiotensin converting enzyme C2 (ACE2) or protease inhibitor 4 (P14), comprising at least one CC polymorphic position as provided in the detailed summary of single CC uncleotide polymorphisms comprising additional 5 and 3 flanking genomic CC sequence; (2) analysing (M1) at least one nucleic acid sample comprising comprising comprising comprising comprising constructing (M2) encoding a protein selected from the group above; (3) constructing (M2) an individual at least two nucleic acid sample comprising comprising grouping at least two nucleic acids (a) identifying (M3) an individual at risk of developing a disorder componesses one or more polymorphic positions in a gene comprises one or more polymorphic positions within a gene encoding a protein selected from the group above; (3) constructing (M2) comprises one or more polymorphic positions within a gene encoding a comprise one or more polymorphic positions within a gene encoding a comprise one or more polymorphic positions within a gene encoding a comprise one or more polymorphic positions within a gene encoding a comprise one position with a known data set. The genes, (M1, M2, M3 and M4) and individual comprising obtaining a nucleic acid sample, determining the construction within a gene encoding a construction within a gene encoding a nucleic acid sample, determining the construction within a gene encoding a nucleic acid sample, determining the construction within a gene encoding a nucleic acid sample, determining at least one polymorphic position, and comparing at least one p
                                         hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory arthritis, cancer, wounds, viral, bacterial or fungal infection, Chronic obstructive pulmonary disease (COPD) and enterocolitis (many other diseases and disorders are listed in the specification). The polynuclectides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence listing but is not referred to anywhere else in the specification
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23-JAN-2001; 2001US-0263678P.
02-MAR-2001; 2001US-0273037P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated nucleic acid from a h
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          autoimmune diseases.
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Swanson BN,
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TSUCHIHASHI Z.
HUI L.
18
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Powell
BP; 1
A; 9 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JR;
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(BDKRB1),
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RESULT 129
ABS60512/c
ID ABS605
             CC encoding aminopeptidase P (XPNEP2); bradykinin receptor B1 (EDRKB1), C1 catchykinin receptor B1 (EDRKB1), C2 catchykinin receptor B2 (EDRKB2), angiotensin converting enzyme C2 (ACE2) or protease inhibitor 4 (P14), comprising at least one C2 (ACE2) or protease inhibitor 4 (P14), comprising at least one C2 (ACE2) or protease inhibitor 4 (P14), comprising at least one C3 (Polymorphic position. Also included are (1) a probe that hybridises to a C4 polymorphic position as provided in the detailed summary of single C5 (P14) (P14), comprising additional 5 and 3 flanking genomic C6 sequence; (2) analysing (M1) at least one nucleic acid sample comprising C7 cobtaining the sequence at one or more individuals and determining the C7 catching a protein selected from the group above; (3) constructing (M2) (P14) and protein games comprising grouping at least two nucleic acids (4) identifying (M3) an individual at risk of developing a disorder C6 (P14) 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein KLK1; bradykinin receptor B2; BDKRB2; gene therapy; angiotensin converting enzyme 2; ACB2; protease inhibitor 4; PI4; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; angioedaema; aneurysm; embolism; thrombosis; coronary artrophy; vascular disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-2000; 2000US-0251015P.
23-JAN-2001; 2001US-0263678P.
02-MAR-2001; 2001US-0273037P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 798; 977pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              autoimmune
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Swanson BN,
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TSUCHIHASHI Z.
HUI L.
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Powell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         isolated nucleic acid from a human
XPNEP2), bradykinin receptor B1 (BD)
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C1NH; ka
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L (BDKRB1),
WH), kallikrein
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RESULT 130
ABL43873/c
ID ABL438
XX ABL438
AX ABL438
AX ABL438
AX Human;
AX Homo E
XX H
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                                                                                     The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. Succeed to the maximum in the specified discrimination Nos. It array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. Succeed to the specified discrimination Nos. (are mixed respectively in each wells of longitudinal and larger and array and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory arthritis, cancer, wounds, viral, bacterial or fungal infection, Chronic obstructive pulmonary disease (COPD) and enterocolitis (many other diseases and disorders are listed in the specification). The polynucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence listing but is not referred to anywhere else in the specification
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and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) sare detected from the amplified products; (h) the clones in the materials.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 23; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arraying genome clones.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human chromosome 1p36-35 PCR primer SEQ ID NO:917.
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(GENO-) GENOTEX YG.
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The invention relates to a method for inhibiting viral replication in a cell or animal, where a heat-shock protein (HSP) is required for replication, comprises administering an inhibitor of the heat shock protein. Also included are a kit for the process comprising the protein and instructions; an isolated nucleic acid that is complementary (antisense) to a sequence that encodes the HSP; a vector containing the antisense molecule; a non-human transgenic mammal containing the antisense molecule; a method for inhibiting viral replication in a cell by administering the antisense molecule; or its argument; a non-human transgenic mammal containing isolated nucleic acid that encodes a viral particle u-binding protein (UBP) or its derivative or fragment; a non-human transgenic mammal containing isolated nucleic acid that encodes an inhibitor of HSP-dependent viral replication; a method for inhibiting viral replication in a cell by administering the inhibitor encoding nucleic acid; and a method for identifying compounds that inhibit HSP-dependent viral replication. The method is used to treat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 C; 7 G; 1 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                       "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.4;
Pred. No. 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IKBR; I-kappa-B-related; NFKBIL2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78;
                                                                                                                                                                                                                                                                                                                                                                                                                                             (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ξ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             All cytidine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ISIS #130489.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases

of inhibitor-kappa B-R,

e.g.,

a heightened

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RESULT 133
AAL61584
ID AAL6158
XX AAL615
XX AAL615
XX AAL615
XX Human;
XX Human;
XX Human;
XX Itagpa
XX It
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to antisense compounds targetted to a nucleic acid comblecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IXBR, I-kappa-B-related, ikappabr, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of increased cytokine expression, or a result of infection (e.g. bacterial, cyiral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between C functions of various members of a biological pathway. They are also cuseful in antisense therapy. The present sequence is an oligonucleotide or targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       associated with immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tumour; prophylaxis; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAL61584;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAL61584 standard; DNA;
                                                                                                                                                                                                                                        WO2003042360-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                              13-NOV-2001; 2001US-00993731
                                                                                                                                 05-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 Similarity
                              SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTGCCTTTGAGAAGAG 790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 BP; 4 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGCCTTGGAGAAGAG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                              PHARM INC
                                                                                                                                   2002WO-US035597.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                   /*tag= b
/mod_base= OTHER
/note= "2'-methoxyethyl (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression of or infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     are
                                                                                                                                                                                                                                                                                                                 /mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93.8%;
                                                                                                                                                                                                                                                                                   _base= OTHER
== "2'-methoxyethyl (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                                                                                ი
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.4;
Pred. No. 1e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      le+02;
                                                                                                                                                                                                                                                                                                                                                                                                      (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                      nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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The invention relates to antisense compounds targetted to a nucleic acid comblecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, CC IKER, I-kappa-B-related, ikappab r, nuclear factor of kappa light color polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to climbibit its expression. Antisense compounds of the invention are useful cfor treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving concreased cytokine expression, or a result of infection (e.g. bacterial, corrected cytokine expression, or a result of infection (e.g. bacterial, corrected cytokine expression) or a result of infection (e.g. bacterial, corrected cytokine expression) or a result of infection of the corrected corrected cytokine expression or a result of infection of the corrected corrected cytokine expression or delay infection, inflammation or tumour corrected corrected cytokine expression of a biological pathway. They are also cuseful in antisense therapy. The present sequence is an oligonucleotide cargetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                 Sequence 20 BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; Page 75; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Watt
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Length 20;
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Matches
                                                                                                                                                                  Query Match
IGF-I oligonucleotide #3781.
                    30-MAR-2001
                                       AAF52821;
                                                         AAF52821 standard;
                                                                                                                                                           Local
                                                                                                                             1252 GGCTGCAGCAACAGCTGGA 1270
                                                                                                                                                 16;
                                                                                                           N
                                                                                                                                                          Similarity
                                                                                                           GGCTGCAGCTCAGCTGCA 20
                                                                                                                                                 Conservative
                   (first entry)
                                                          DNA;
                                                                                                                                                          0.6%;
                                                           15
                                                          ВP
                                                                                                                                                 0
                                                                                                                                                          Score 14.2;
Pred. No. 1
                                                                                                                                                 ed. No. 1.1e+02;
Mismatches 3
                                                                                                                                                                     DB 1;
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片 S

RESULT 134
AAF558
ID AAF558
XX AAF558
XX AAF558
XX AAF558
XX AAF558
XX AAF558
XX AAF566
XX GF0Wh
KW IGF bi
KW IGF bi Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; seleroderma; wart; skin cancer; sclerotic disease; keratosis; neoplasia; shopting disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss. sapiens.

WO200078341-A1

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693

21-JUN-1999; 99US-0140345P

(MURD-) MURDOCH CHILDRENS RES INST

WPI; 2001-041421/05 С С

Werther

GA,

Edmondson

Ameliorating the effects of. ĝ disorder, e.g. psoriasis, by administering

treatment

that

Example

6; Page 34; 201pp; English

PERKERA SOCIO SOCI

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RESULT 135
AAF451
XX AAF45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; opthhalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFP-2; IGFPB-3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; hyperneovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 UV (ultra-violet) inhibits or reduce inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Page 85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF45187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15
                                                             Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WC200078341-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUN-2000; 2000WO-AU000693
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                                                                                                                                                                                                                                                                                                                                                                                    21-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention relates to a method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1232 GCATGTGCTGGCAG 1245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                       MURDOCH CHILDRENS RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GCATGTGCTGGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   reduces
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                         Werther
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cc skin disorders. The method comprises contacting the effects of cantisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of cinhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense CC oligonucleotides of the present invention (see AAF45151 and AAF45153-CC (CC F45161). The method is useful for ameliorating the effects of psoriasis, cc inchiposis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC neoplasias, scleroderma, warts, benign growths, cancers of the skin, and continuous condition of the retina, cc hyperneovascular condition such as a neovascular condition of the retina, cc brain or skin, growth factor-mediated malignancies, other sclerotic cc vessels or any other hyperplasia
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                                                                                                                                                   WPI; 2001-041421/05
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                                                                                                        Ameliorating the effects of a disorder, UV (ultra-violet) treatment (optional)
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The present Example 6; Page

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RESULT 137
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AAF48301
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                                                                                                                                                                                                                                                                                                Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200078341-A1
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The present invention relates to a method for ameliorating the effect skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an

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oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psorias ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosi

The present invention relates to a method for ameliorating the effect skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide of the present invention (see ARP45151 and ARP45153-P45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhosa, kelodids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
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RESULT 141
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                                                                                                                                                                                                                       The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcriptio factor gene, IRF-2 and/or the CATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production erythropoietin, granulocyte colony stimulating factor protein and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor prointerferon alpha and erythropoietin.
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                                                                                   100.0%;
                                                                                                         0.6%;
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Mismatches
                                                           Mismatches
                                                                                DB 1;
81;
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                                                        0
                                                                                                         Length 17
                                                                                                                                                                                                                                                                                                                                           n of repressor genes
the GATA transcription
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RESULT 142
AAF07203/c
ID AAF072
XX AAF072
XX AAF072
XX AAF072
XX AAF072
XX Hammer
XX Ribozy
KW Interf
XX Homo s
XX Homo s
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YN W02000
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Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ribozyme; erythropoietin; interferon alpha; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 0 A;
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                                                                                                                                              Ribozyme; erythropoietin;
                                                                                                                                                                    Hammerhead ribozyme substrate #3460.
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                                    11-APR-2000; 2000WO-US009721.
                                                                                   WO200061729-A2
                                                                                                          Homo sapiens
                                                                                                                                                                                                                    AAF07203;
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              12-APR-1999;
                                                                                                                                 interferon alpha;
                                                                                                                                                                                             16-FEB-2001
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            99US-0129390P
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                                                                                                                                                                                             entry)
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                                                                                                                                              granulocyte colony stimulating
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Pred. No.
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RESULT 143
ABK023
XX ABK023
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KW Human;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NGG; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; NGL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy induced neuropathy; amyotrophic lateral sclerosis; ALS; parkingon's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                11-FEB-2000;
28-FEB-2000;
06-MAR-2000;
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                                                     (RIBO-)
(BLAT/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
Synthetic.
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                                                                                RIBOZYME PHARM INC.
BLATT L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             alpha
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                                                                                                                                                                2000US-0181797P.
2000US-0185516P.
2000US-0187128P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA;
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MCSWIGGEN J CHOWRIRA B M

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ABKO2096

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CC possessing an NCH motti) and NGN triplet), a zinzyme (cleaving RNA with a YGY motti). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2+.

CC processing and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more thoughtin's lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-cc leukaemia, HIV (human immunodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), immunocyrcoma (IMC), small B-cell lymphocyric lymphoma, cc immune thrombocyrcoparia, and inflammatory arthropathy. The NOGO-crested with a cell to reduce NGO gene in the ccl arigetting nucleic acid is used to cleave RNA of the NGO gene in the ccl acid may be contacted with a cell to reduce NGO activity of the CC largetting nucleic acid may be used to cleave RNA to the NGO activity of the CC cell and treat a patient having a condition associated with the level of therapies. In particular, the NGO-targetting nucleic acid may be used to cleave contacted with a cell to reduce NGO activity of the coll and treat a patient having a condition associated with the level of CC patiency induced neuropathy, amyotrophic lateral sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), chemotherapy induced to the modulation of NGO expression. T
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Best Local S
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Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic, neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantie-cell lymphoma; NCL; immunocytoma; IMC; immunocytoma; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                      12-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1884 GAGGAGGACGAGGA 1897
                                                                                                                                                                                                                                                                                                                                                                   NOGO
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Matches Query Match Best Local

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DNAzyme molecule

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down conclusive states expression of a neurite growth inhibitor gene (NOGO). The conclusion and the enzymetic nucleic acids (e.g. a ribozyme or a condition and the enzymetic nucleic acids (e.g. a ribozyme or a condition and the enzymetic nucleic acids (e.g. a ribozyme or a condition and the enzymetic nucleic acid cleaving a new NNA molecule consessing an NCH motif), a G-cleaver (cleaving RNA with a NNA motif) proceed an amberzyme (cleaving RNA with a NNA triplet), a zinzyme (cleaving RNA with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therepies. In particular, the CD20 targetting nucleic acid may be used to cleave RNA condition associated with the level conformation (NCL), immunocytoma (NMC), small B-cell lymphocytic lymphoma, leukaemia, Hr (human immunodeficiency virus) associated NHL, lymphocytic conformation associated with the cell cardid may be contacted with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably Mg^2+. Furthermore, the cell individual new particular, the NOGO-cent cardid with the level of conformation associated with the level of conformation acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of conformation and conformation and condition associated with th
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28-FEB-2000;
06-MAR-2000;
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(BLAT/)
(MCSW/)
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Parkinson's disease; ataxia; Huntington's disease;
Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease
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BLATT L.
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RESULT 145
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XX Claim
The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down CC regulates expression of a neurite growth inhibitor gene (NGGO). The CC nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CC DNRzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule CC possessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) pr con amberzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20 targetting nucleic acid is used to cleave RNA CC with a YGY motif). The CD20 targetting nucleic acid is used to cleave RNA CC price of a divalent cation that is preferably Mg^2+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHI, lymphoma, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHI, mantle-cell lymphoma (MCI), immunocytoma (IMC), small B-cell lymphocytic lymphoma, in the condition and inflammatory arthropathy. The NGGO-
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28-FEB-2000;
06-MAR-2000;
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(BLAT/)
(MCSW/)
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Synthetic.
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) BLATT L.
) MCSWIGGEN J.
) CHOWRIRA B M.
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and

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

2002-179446/23

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RESULT 146
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Best Local Similarity
Matches 14; Conserv
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30-JAN-2001;
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30-JAN-2001;
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27-SEP-2000;
04-OCT-2000;
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Pred. No.
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RESULT 147
ABNO2623
ID ABNO26
XX ABNO26
XX Human
XX Human;
KW Human;
KW Muscle
KW Skelet
XX WO2001
XX WO2001
XX WO2001
XX O6-DEC
XX O6-D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 cc nucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to cc provide initial substrates for the recombinant engineering of hGDMLP-1 cc protein variants having desired phenotypic improvements, and for cc expressing the proteins. The hGDMLP-1 proteins or polypeptides may be closed as immunogens to raise antibodies that specifically recognise hGDMLP consider a second proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser descrption ionisation, as continuously the proteins and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The collision of disorder associated with the expression of hGDMLP-1, in particular heart can disorder associated with the expression of hGDMLP-1, in particular heart can disorder associated with the expression of hGDMLP-1, in particular heart can be guence disorders. hGDMLP-1 is localised to chromosome 2.

The present sequence in the exemplification of the present invention. N.B. complete data for this patent did not form part of the printed continuous contraction, but was obtained in electronic format directly from MIPO cat figure in the screening of the contraction of the printed contraction of the printed contraction.
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                         26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    skeletal muscle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human GDMLP-1 17-mer scanning
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14; Conserv
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                           2001WO-US000667
2001WO-US000663
2001WO-US000663
2001WO-US000664
2001WO-US000666
2001WO-US000667
2001WO-US000667
2001WO-US000667
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2001WO-US000667
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larity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                   2000US-0207456P.
2000US-0234687P.
2000US-0236359P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14;
Pred. No.
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81;
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Sequence 17

BP; 5

A; 4 C; 6

<u>.</u>

2 T; 0

다; 0

Other;

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The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 cC nucleic acids can be used as probes to detect, characterise and quantify comproved initial substrates for the recombinant engineering of hGDMLP-1 cC protein variants having desired phenotypic improvements, and for cexpressing the proteins. The hGDMLP-1 proteins or polypeptides may be compressed in the proteins. The hGDMLP-1 proteins or polypeptides may be compressed in the proteins as standards in assays used to determine the concentration capture probes for surface-enhanced laser described initiation, as capture probes for surface-enhanced laser described energy in hGDMLP-1 capture probes for surface-enhanced laser described energy in hGDMLP-1 composition, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a cdisorder associated with the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present invention. N.B. The present sequence in the exemplification of the present invention. N.B. ChGDMLP-1 sequence data for this pattent did not form part of the printed composition, but was obtained in electronic format directly from WIPO at figure 1 and 1 and 1 and 1 particular sequence control of the printed contr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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Ş 밁 Query Match Best Local S Matches 14 1840 l Similarity 14; Conserv TCTCAGAGAGCGAG 1853 TCTCAGAGAGCGAG Conservative 0; 0.6%; 17 Score 14; Pred. No. Mismatches B1; ۲, <u>,,</u> Length 17 Indels 0, Gaps 0

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RESULT 148
ABK17922/c
ID ABK17922;
XX
AC ABK17922;
XX
C ABK17922;
XX
DE Human ERG hammerhead ribozyme, of the company of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human ERG hammerhead ribozyme target sequence,
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Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopath vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopath; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulga; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; sturge weber syndrome; kippel-Trenaunay-Weber syndrome; kippel-Trenaunay-Weber syndrome; leukaemia; sturge weber syndrome; kippel-Trenaunay-Weber sy leukaemia; osteoporosis; DNAzyme; virucide; osteopathic; vulgaris;

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RESULT 149
ABK17921/C
ID ABK179
XX ABK179
XX O9-APR
XX U9-APR
XX Human;
XW Uphrha
XW Vulne;
XW THOWAGE
XW THOWAGE
XW THOWAGE
XW THOWAGE
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XW ABG101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a nucleic acid molecule (I) which down regulates coxpression of an Ets-related gene (ERG), (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour anglogenesis, diabetto retinopathy, macular degeneration, convascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or cell, by contacting the cell with (I). (I) is useful for reducing ERG activity in a cell, by contacting (I) with RNA, in the presence of a divalent cation such as my2+. (I) is useful for reducing ERG activity in a cell, so the expression of ERG, and as diagnostic tool to the presence of ERG RNA in a cell. (I) is useful for specifically cation such as material such as a diagnostic tool to the presence of ERG RNA in a cell. (I) is useful for specifically cation such as mare homology with ERG gene or ERG fusion genes. CRI and mutations within ERG gene or ERG fusion genes. CRI activity in a cell caids, including antisense and centry and contacting caids and cells or to detect the presence of a caid molecules which regulate expression of ERG, and cells or to detect the presence of activity in a cell caids, including antisense and cell cells or to detect the presence of the invention
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Best Local Sim
Matches, 14;
                  Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsorialic; virucide; osteopathi vulnerary; cancer; lymphoma; Ewing's Sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgar angiofibroma of tuberous sclerosis; port-wine stain; wound healing; sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; soler_Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozym
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                                                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                                                                            ABK17921
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14
                                                                                                                                                                                                                      ERG hammerhead ribozyme target sequence, Seq
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GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                            standard;
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                                                                                                                                                                                                                                                                                                                                                          RNA;
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                                                                                                                                                           osteopathic;
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                                                                                           vulgaris;
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RESULT 150
ADB42705/c
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SXBXBBXBXB
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Best Local
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                              Tumour suppression/reversion associated nucleotide #3028
                                                                   18-DEC-2003
04-DEC-2003
                                                                                                                                                           ADB42705 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 69; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jarvis T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                        ADB42705;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                     1880 GCTGGAGGAGGACG 1893
                                                                                                                                                                                                                                                                                                                        al Similarity
14; Conserv
                                                                                                                                                                                                                                                   片
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Von Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                             BP; 2 A; 8
                                                                                                                                                                                                                                                                                                                          Conservative
                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                            0.6%;
                                                                                                                                                                                                                                                                                                                                                                                             C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                           Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen JA,
                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                             , 1B
                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mclaughlin
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                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     down regulates
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                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  fur in figure 1 of ERG,
                                                                                                                                                                                                                                                                                                                            Gaps
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0

cytostatic;

antiviral;

neuroprotective;

nootropic; neuroleptic;

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ART44257
ID AAT442
XX AAT442
AC AAT442
XX 22-JUI
XX 22-JUI
XX XX 22-JUI
XX XX Antise
KW Antise
XX Nuclee
XX XX Synthe
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                                                                                                                                                                                                                                                                                                  ઠ
                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to the isolation of 6327 nucleotide sequences, a fragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of nucleotides involved in tunour suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing them and cells containing the vectors), the encoded polypeptides and antibodies (Alb) against the polypeptide are useful for prevention and/or treatment of viral infections or diseases characterized by development of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or cell degeneration (e.g. Alzheimer's disease or schizophrenia).
Analysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can also be used to screen for their specific interactive molecules, potentially useful for treating diseases associated with abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 386; 771pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful e.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-441574/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-2001; 2001FR-00011981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-2002; 2002WO-IB004219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003040369-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; probe; tumour suppression; tumour reversion; apoptosis;
virus resistance; transgenic animals; Alzheimer's disease; schi
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 2 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression of the nucleotides.
                                                Antisense therapy; herpes simplex virus; HSV-1; guanosine; nuclease resistance; stability; ss.
                                                                                                                                          22-JUL-1997
                                                                                                                                                                                                            AAT44257 standard;
                                                                                                   HSV-1 antisense component of capped oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid encoding human prostate membrane-specific antigen, ful e.g. for treatment of tumors and viral infection, also related ypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                  1018 GCCAGCACTCCCAG 1031
                                                                                                                                                                                                                                                                                                  17
                                                                                                                                                                                                                                                                                                                                                                      14;
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                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amson R,
                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGINES
                                                                                                                                                                                                              DNA; 17
                                                                                                                                       entry!
                                                                                                                                                                                                                                                                                                                                                                                       0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tuijnder M;
                                                                                                                                                                                                              뜅
                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                       Score 14; DB 1; Length 17; Pred. No. 81;
                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    schizophrenia;
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RESULT 152
AAX3390
XX AAX339
XX AAX339
XX AAX339
XX JO-JUN
XX Gene e
KW viral e
KW cell-c
XX Synthe
OS Human
XX Synthe
OS Human
XX 12-MAH
PH 12-MAH
PR 13-MAH
PR 13-MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RANGE SERVICE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.5%;
Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 4 A; 1 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ten- to 40-mer oligonucleotides which have a cap of 1-10 (esp. 4) G residues on at least one end are provided; if caps are present at both ends, they can be of the same or different lengths. A cap sequence increases nuclease resistance of the oligonucleotide and also increases cell penetration. The present sequence is that of a preferred oligonucleotide, directed against HSV-1 sequences, which can be capped
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligo:nucleotide(s) with series of G residues at at least one end have increased stability against nuclease and cell penetration, - are particanti:sense sequences for treating and diagnosing cancer, viral diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DE19502912-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell-cell adhesion receptor; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene expression inhibitor; probe; nucleic acid detection; growth factor;
viral infection; therapy; HSV-1; cancer; restenosis; integrin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSV-1 gene expression inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX33904 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peyman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FARH ) HOECHST AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for use in antiviral therapy
                                                                                                                                                                                                                                                                                                                   13-MAR-1995;
24-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AU9648028-A
                                                                                 WPI; 1996-455932/46
                                                                                                                                                                                                                                                                                                                                                                                                                                      12-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human herpesvirus
                                                                                                                                                                                                                                          (FARH ) HOECHST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1775 GGAGGAGGCGGAGGAGG 1791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Uhlmann E;
                                                                                                                                                         Uhlmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95DE-01002912.
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                                                                                                                                                                                                                                                                                                                       95DE-01008923.
95DE-01043865.
                                                                                                                                                                                                                                                                                                                                                                                                                                               96AU-00048028.
                                                                                                                                                                                                                                          AG.
                                                                                                                                                         Ħ
                                                                                                                                                         Breipohl
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Pred. No. 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                  Wallmeier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ٥,
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            of gene
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New phosphono:mono:ester oligo:nucleotide analogues

inhibitors

FX8X555555555555XX

밁 S

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RESULT 153
AAT81044/c
ID AAT810
XX AAT810
XX AAT810
XX AAT810
XX Enzyma
KW Enzyma
KW Enzyma
KW COFONA
XX Enzyma
XX HOMO S
XX IS-MAY
XX Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression (antisense oligonucleotides, ribozymes, sense oligonucleotides and triplex-forming oligonucleotides), as probes for the detection of nucleic acids, and as auxiliaries in molecular biology. As gene expression inhibitors they may be used for treating viral infections (especially where the virus is HSV-1, HSV-2, an influenza virus, VSV, hepatitis B or papilloma virus), cancer, restenosis, medical conditions mediated by integrins or cell-cell adhesion receptors, and medical conditions induced by growth factors (especially TNF-alpha)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents an inhibitor of HSV-1 gene expression, and example of an oligonuclectide analogue of the invention. The oligonuclectide analogues of the invention are used as inhibitors of
The present sequence represents the preferred target sequence for an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes the c-myb sequence and can be used to prevent smooth muscle cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enzymatic nucleic acid; hammerhead; ribozyme; cleavage; human; smooth muscle cell; hyperproliferation; restenosis; cancer: c-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression
                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-MAY-1994;
13-JAN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                         produced by e.g. c-myb,
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                              The present sequence represents the preferred target sequence for an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes cleave the c-myb sequence and can be used to prevent smooth muscle cell hyperproliferation in restenosis, especially after coronary angioplasty,
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                       the c-myb sequence hyperproliferation and in cancers
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                                                                                                                                                                                                                                                      Stinchcomb DT,
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13-JAN-1995;
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RESULT 155
AAT81047/C
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RESULT 156
AAT81045/c
ID AAT810
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DT 26-SBE
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Best Local S
Matches 15
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13-JAN-1995;
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smooth muscle cell; hyperproliferation; restenosis; cancer; c-myb;
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                         Enzymatic nucleic acid; hammerhead; ribozyme; cleavage; human;
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smooth muscle
                                                                                Human c-myb hammerhead ribozyme target sequence (nt.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                    (first
cell; hyperproliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94US-00245466.
95US-00373124.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHARM INC.
                                                                                                                                                                                                                                                 RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.5%;
                                                                                                                                                                                                                                                 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 13.8;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cleave RNA produced by e.g. c-myb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                   position 28)
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RESULT 157
AAX2421
ID AAX242
XX AAX242
XX O1-JUI
DT 01-JUI
DT 01-JUI
DX Phosph
XX Phosph
XX Phosph
XX Integr
XX Integr
XX DE195(
XX DE19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents the preferred target sequence for an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the maNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes cleave the c-myb sequence and can be used to prevent smooth muscle cell hyperproliferation in restenosis, especially after coronary angioplasty.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 coronary angioplasty; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the c-myb sequence hyperproliferation and in cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New enzymatic nucleic acid molecules - for treating restenosis or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1996-010927/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stinchcomb DT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9531541-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX24211 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 64; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-MAY-1994;
13-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-NOV-1995.
                                                                                                                                                                                                                                                                                                                      Phosphonomonoester analogue; inhibitor; antisense; cancer; ribozyme; diagnostic agent; detection; treatment; disease; integrin; cell-cell adhesion receptor; TNF-alpha; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                     Phosphonomonoester
                                                           13-MAR-1995;
                                                                                                               13-MAR-1995;
                                                                                                                                                                  19-SEP-1996
                                                                                                                                                                                                                        DE19508923-A1
     (FARH ) HOECHST AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGGAGGAGGCGGAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 0 A; 10 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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95US-00373124
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                                                           95DE-01008923
                                                                                                                  95DE-01008923
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                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotide analogue 28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 13.8;
Pred. No. 88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cleave RNA produced by e.g. c-myb,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Length
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                                                                                                                                                                                                                                                                                                                                                                 virus;
                                                                                                                                                                                                                                                                                                                                                                                              restenosis;
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RESULT 158
AAT09036/c
ID AAT090
XX AAT090
AC AAT090
XX 28-AUG
DD Arabid
XX EIN2;
KW Ethyle
XX W09535
XX W09535
XX I7-JUN
XX I7-JUN
XX I7-JUN
XX Phan
Ecker
XX WPI; 1
XX Plante
PT Plante
P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel phosphonomonoester oligonucleotide analogues which act as inhibitors of gene expression (as sense/antisense, ribozyme or triplex-forming molecules), useful as diagnostic agents (i.e. probes for detecting nucleic acid) or seful as diagnostic agents (i.e. probes for detecting nucleic acid) or of or triplex-forming molecules (i.e. probes for detecting nucleic acid) or of or triplex-forming molecules (i.e. probes for detecting nucleic acid) or of or triplex-forming molecules (i.e. probes for detecting nucleic acid) or of or triplex-forming molecules (i.e. probes for detecting nucleic acid) or of or other probes for detecting nucleic acid) or of or other probes for detecting nucleic acid) or of or other probes for detecting nucleic acid) or of or other probes for detecting nucleic acid) or other probes for detecting nucleic acid or oth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New oligo:nucleotide analogues contg. phospho:mono:ester bridges - therapeutic inhibition of gene expression, e.g. in cancer or viral infection, with good specificity and in vivo stability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             viruses, influenced by integrins or cell-cell adhesion receptors, induced by factors such as TNF-alpha, or cancer or restenosis. The products of the invention satisfy the requirements of good in-vivo stability; ability to cross cellular and nuclear membranes, and specific binding to target nucleic acid better than known oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 23; 36pp; German
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          The present sequence is a primer for the A. thaliana EIN2 (ethylene insensitive) locus. When transformed into plants EIN2 genomic DNA, or CDNA sequences (obtd. from the EIN2 locus) confer disease tolerance and ethylene insensitivity, with minimal injury or reduction in the harvest yield of saleable material. The plants with disease tolerance may have extensive levels of infection, but little necrosis and few or no lesions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plant sequences for ethylene
confer disease tolerance and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ethylene insensitivity; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1775 GGAGGAGGCGGAGGAGG 1791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ethylene insensitive; transformed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rothenberg
                                                                                                                                                                                                                                                                                                                                      Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
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                                                                                                                                                                                                                                                                                                                                           144pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EIN2 (ethylene insensitive)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              insensitive loci and hook-less 1 allele(s) ethylene insensitivity when transformed in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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11-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  They may also have reduced necrotic and water soaking chlorophyll loss may be virtually absent
                      The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRWA encoding 1 or more receptors of vascular endothelial growth factor (NGGP). A patient preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                       Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
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tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
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    Sequence
                                                                                                                                                                  Claim 4;
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Matches 10
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                                                                                                                                                                                                                                                                                                                                                                                                               stability - useful for treating e.g. turbeumatoid arthritis, etc., in a human
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11-JAN-1996;
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                                                              RNA; 17
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Human NOGO Amberzyme
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cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme MNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHI; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHI; mantle-cell lymphoma; MCI; immunocytoma; INC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease; Human; ss; antisense therapy; cytostatic; antiinflammatory; disease; muscular dystrophy; neurodegenerative disease hammerhead ribozyme; haemostatic;

Homo sapiens. Synthetic.

WO200159103-A2

16-AUG-2001.

09-FEB-2001; 2001WO-US004273

11-FEB-2000; 28-FEB-2000; 06-MAR-2000; ; 2000US-0181797P. ; 2000US-0185516P. ; 2000US-0187128P.

(RIBO-) (BLAT/) (MCSW/)) RIBOZYME PHARM :
) BLATT L.
) MCSWIGGEN J.

Mcswiggen J, Chowrira

CHOWRIRA B

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, central nervous system injury.

Claim 131; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates corpression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a comparison of a neurite growth inhibitor gene (NGGO). The comparison of a nucleic acids (e.g. a ribozyme or a comparison of a nucleic acids (e.g. a ribozyme or a comparison of cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) processor of the cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a CD20) trate is preferably Mg^2 + the comparison of the collague RNA of CD20 in the presence of a divalent cation that is preferably Mg^2 + conferency it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of targetting nucleic acid may be used to treat tymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHL, lymphocytic lymphoma, low-grade or follicular NHL, lymphoma, low-grade lymphoma, low-grade lymphoma, low-grade lymphoma, low-grade lymph

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RESULT 162
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                                                                                                                                                                                                                                                                                       Nucleic acid molecules, e.g., enzymatic nucleic acids and constructs, which down regulate expression of a CD20 gene growth inhibitor gene useful for treating, e.g., lymphoma,
  The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) pr an amberzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA)
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                                                                                                                                                                                                                                                                      central nervous
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                                                                                                                                                                                                                                                                   system injury.
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Pred. No. 88
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or neurite
a, leukemia,
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CC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA CO of CD20 in the presence of a divalent cation that is preferably Mg^2^+. CC Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level CC of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-CC (Hodgkin's lymphoma (NHL), bulky low-grade or follicular MHL, lymphocytic (leukaemia, HCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immunodeficiency virus) associated MHL, mantle-cell (lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, climant trombccytopaenia, and inflammatory arthropathy. The NCGO-CC (leukaemia, HCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, climant trombccytopaenia, and inflammatory arthropathy. The NCGO-CC (leukaemia, HCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, climant trombccytopaenia, and inflammatory arthropathy. The NCGO-CC (leukaemia, HCC) acid may be contacted with the CC (leukaemia, HCC) acid may be contacted with a cell to reduce NCGO activity of the CC (leukaemia) be contacted with a cell to reduce NCGO activity of the CC (leukaemia) be contacted with a cell to reduce NCGO activity of the CC (leukaemia) be contacted with the level of the contacted with the level of the contacted with the level of the seasociated with the level of the seasociated with the level of coll may be contacted with the level of the seasociated with the level of the seasociated with the level of coll may be contacted wi
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                                                                              amberzyme molecule of the invention
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Sequence 17 BP; 8 A; 0 C; 9 G; 0 T; 0 U; 0 Other;

Matches Query Match Best Local S 1764 GAAGATGAGGAGGAGGA 1780 15 15 Similarity Conservative 88.2%; 17 0 Score Pred. Mismatches ŏ 13.8; DB <u>,</u> ν •• Length 17; Indels 0 Gaps 0

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RESULT 163 ABK02462 PRESENTATION OF STREET PRODUCTION OF STREET PRODUCT 12-MAR-2002 ABK02462; ABK02462 standard; RNA; 17 (first entry) BP.

Human NOGO

Amberzyme #134.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NGG0; hammerhead ribozyme; NAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; b-cell lymphoma; non-Hodgkin's lymphoma; NHI; hymphocytic leukaemia; human immunodeficiency virus; HIV associated NHI; mantle-cell lymphoma; NGL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy induced neuropathy; amyotrophic lateral sclerosis; ALS; parkingon's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease

Synthetic Homo sapiens

WC200159103-A2

16-AUG-2001.

11-FEB-2000; 2000US-0181797F 28-FEB-2000; 2000US-0185516F 06-MAR-2000; 2000US-0187128F 09-FEB-2001; 2001WO-US004273.

(RIBO-) RIBOZYME PHARM INC

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                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; Lnozyme, G-Cleaver; amberzyme; zinzyme; Lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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(MCSW/)
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) MCSWIGGEN J.
) CHOWRIRA B M.
                                                               ss; antisense therapy; cytostatic; antiinflammatory; haemostatic;
oprotective; nootropic; neuroprotective; antiparkinsonian;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acid molecules, e.g.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                  (first entry
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CC expression of a CD20 gene and a nucleic acid molecule which down communication and a CD20 gene and a nucleic acids of a CD20. The comparison of a neurite growth inhibitor gene (NOGO). The comparison of a neurite growth inhibitor gene (NOGO). The comparison of a comparison of a comparison of a comparison of a comparison of comparison 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 88; Page 131; 200pp; English
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BLATT L.
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amberzyme
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     of the
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Best Local Similarity

Sequence

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13.8; No. 88; 0 U; 0 DB 1;

Length

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28-FEB-2000;
06-MAR-2000;
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-tazgetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2^+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise these of one or more theraples. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurity growth inhibitor gene useful for treating, e.g., lymphoma, leukemia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       central nervous system injury.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-)
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BLATT L.
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CHOWRIRA B 1
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2000US-0185516P.
2000US-0187128P.
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leukemia,
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RESULT 166
ABK023
XX ABK023
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XX Human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                          11-FEB-2000; 2000US-0181797F
28-FEB-2000; 2000US-0185516F
06-MAR-2000; 2000US-0187128F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence is an inozyme of the invention
                                                                             Blatt L,
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                                                                                                                                                                                                                                                                                         (RIBO-)
                                                                                                                                                                                                                                              (BLAT/)
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                                                                                                                                                                                                  RIBOZYME PHARM INC
BLATT L.
MCSWIGGEN J.
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                                                                                                                                                                    CHOWRIRA B
                                                                                  Mcswiggen J,
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Pred. No. 8
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WPI; 2001-607195/69

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The invention relates to a nucleic acid molecule which down regulates conservession of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The CC nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CC DNazyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an RNF motif), a G-cleaver (cleaving RNA with a NYM motif) processing an environment of color of color. The treatment may further comprise the use of one or more contacted (MHL) bulky low-grade or follicular MHL, lymphoma, leukaemia, HrV (human immunodeficiency virus) associated WHL, mantle-cell lymphoma (MCL), immunocytcoma (IMC), small B-cell lymphoma, lymphocytic concessence of a divalent cation that is preferably. The MCGO color of co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nervous system injury.
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Sequence 17 BP; 8 A; 0 C; 9 o G T; 0 ď; 0 Other;

5 Query Match Best Local S Matches 15 1765 l Similarity AAGATGAGGAGGAGGAG 0.5%; nilarity 88.2%; Conservative 1781 0; Score 13.8; Di Pred. No. 88; 0; Mismatches DB 1; 2: Length 17; Indels 0 Gaps 0

RESULT 167 ABA78082/c Н 17

24-JAN-2002 (first entry)

ABA78082

standard;

DNA;

ВP

BRCA1 mutation correcting oligonucleotide SEQ ID

NO:

928.

BRCA1 mutation 24-JAN-2002

correcting oligonucleotide SEQ

IJ

NO:

(first

entry)

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2h; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic;

Homo sapiens

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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the coligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, trethoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A crethoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDXN2A), APC, Pactor V, Factor VIII, Factor IX, haemoglobin alpha locus 2 (HBA1), WHH1, MSH2, MSH6, applipaprotein E (APOB), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APC), presenilin-1 (PSEN) and compresention adenosine deaminase deficiency, cystic fibrosis, thempshilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting coligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-MAR-2000;
27-MAR-2000;
01-JUN-2000;
30-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
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                                                                                                                                                                                                                                                                                                                           Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EB,
                                                                                                                                                                                                                                                                                                                           Page 100; 294pp; English
                                                                                                                                                                                                                                                                                                                                                                               cystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; 2000US-0192176P.
; 2000US-0192179P.
; 2000US-0208538P.
; 2000US-0244989P.
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                                                                                                                                                                                                                                                                                                                                                                             for targeted alterations of genetic sequences fibrosis, comprises at least one mismatch and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rice
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and chemical

Sequence 17 BP; 3 A; 8 C; 2 G; 4 T; 0 ς, 0 Other

oligonucleotides of the invention

Query Match Best Local S Matches 15 15; Similarity Conservative 0.5%; 0 Score 13.8; Pred. No. 8 Mismatches 88; BB <u>ب</u> 2; Length 17; Indels 0 Gaps 0,

片 Ś 1264 AGCTGGAAGAGGCTGAG 1280 17

RESULT 168 ABA78081 standard; DNA; 17 ABA78081;

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor XI; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UCT1; syndrome; APP; PSRM1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCAI, BRCAZ, CFTR, cyclin-dependent kinase inhibitor 2A retinoblastoma.
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27-MAR-2000; 2000US-0192179P.
01-JUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
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modification.
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                                                                         WO200192524-A2
                                                                                                                                                                               Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                       Human GDMLP-1 17-mer scanning
                                                                                                                                                                                                                                                                                                                                             29-MAY-2002
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RESULT 170
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DE Human

ABN02601 standard; DNA; 17

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29-MAY-2002 ABN02601;

(first entry)

Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID

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                                                                                                                                                           therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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21-SEP-2000;
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The present invention describes a human genome-derived myosin-like CC protein 1 (hcDMLP-1). The protein and polynucleotide sequences of hGDMLP-CC 1 can be used in gene therapy and vaccine production. The hGDMLP-1 curleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 curved in variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 capture probes for surface-enhanced laser description deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a construction and in the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The present sequence data for this pattent did not form part of the printed considered in the screening and the form part of the printed considered in the screening of the present invention, but was obtained in electronic format directly from WIPO at fire present did not form part of the printed considered proteins and the company to the present did not form part of the printed considered proteins and the company to the present did not form part of the printed considered proteins and printed directly from WIPO at fire present did not form part of the printed considered proteins and the considered proteins and the considered proteins and the considered proteins and the considered proteins and th
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The present invention describes a human genome-derived myosin-like protein 1 (hgDMLP-1). The protein and polynucleotide sequences of hgDMLP-1 color in gene therapy and vaccine production. The hgDMLP-1 color incleic acids can be used as probes to detect, characterise and quantify complete incleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hgDMLP-1 color variants having desired phenotypic improvements, and for expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hgDMLP proteins, as specific biomolecule capture probes for surface-enhance laser description ionisation, as the capture probes for surface-enhance laser description ionisation, as complement in patients having specific deficiency in hgDMLP-1 colored associated with the expression of hgDMLP-1 may be used for diagnosing a disorder associated with the expression of hgDMLP-1 in particular heart cand skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22. The present sequence in the exemplification of the present invention. N.B. The sequence of the spatial did not form part of the printed consideration, but was obtained in electronic format directly from WIPO specification, but was obtained in electronic format directly from WIPO consideration.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

2002-179446/23.

Disclosure; SEQ ID NO 8082; 214pp; English.

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RESULT 171
ABNO8090
AD ABNO80
AX BABNO80
AX BABNO80
AX BABNO80
AX BABNO80
AX BABNO80
AX ABNO80
AX ABNO80
AX ABNO80
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30-JAN-2001;
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27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JAN-2001;
30-JAN-2001;
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2000US-0236359P.
2000US-0236359P.
2000US-0236359P.
2001WO-US000661.
2001WO-US000662.
2001WO-US000663.
2001WO-US000665.
2001WO-US000665.
2001WO-US000665.
2001WO-US000669.
2001WO-US000669.
2001WO-US000669.
2001WO-US000669.
2001WO-US000669.
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ftp.wipo.int/pub/published_pct_sequence

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RESULT 172
ABNO7862/c
ID ABNO7862;
XX ABNO7862;
XX ABNO7862;
XX ABNO7862;
XX ABNO7862;
XX Human GDMI
XX Human; gen
KW muscle; m)
KW Skeletal r
XX Homo sapis
XX WO2001925;
XX WO2001925;
XX WO2001925;
XX WO2001925;
XX O6-DEC-20
XX 25-MAY-20
PR 21-SEP-20
PR 21-SEP-20
PR 21-SEP-20
PR 21-SEP-20
PR 30-JAN-20

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Best Local Similarity 88.3
Matches 15; Conservative
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
The present invention describes a human genome-derived myosin-like protein 1 (homMLP-1). The protein and polynucleotide sequences of homMLP-1 can be used in gene therapy and vaccine production. The homMLP-1 nucleic acids can be used as probes to detect, characterise and quantify homolule-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of homMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The homolule-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise homolule-1 proteins, as standards in assays used to determine the concentration
                                                                                                                                                                                                                                                                                                                                                   New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAY-2001; 2001WO-US016981
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2000US-023458PP.
2000US-0236359P.
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2001WO-US000661.
2001WO-US000662.
2001WO-US000664.
2001WO-US000665.
2001WO-US000667.
2001WO-US000668.
2001WO-US000668.
2001WO-US000668.
2001WO-US000668.
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Pred. No. 88;
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RESULT 173
ABN10746/c
ID ABN107
XX ABN107
XX ABN107
XX ABN107
XX ABN107
XX Human;
XX Human;
XX Human;
XX Human;
XX Homo s
XX WO2001
XX Belet
XX WO2001
XX 26-MAY
XX 26-MAY
XX 26-MAY
PR 21-SEP
PR 27-SEP
PR 27-SEP
PR 27-SEP
PR 30-JAN
PR 30
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Best Local &
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30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001WS-0266860P.
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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2001WO-US000661.
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2000US-0234687P.
2000US-0236359P.
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Pred. No. 88;
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New polypeptide, for raising antibodies that recognize hGDMLP-1 protor as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

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RESULT 174
ABN1071A8/c
ID ABN107
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XX Human;
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21-SEP-2000;
27-SEP-2000;
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2000US-0234687P.
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Correction 1 (hopMilp-1). The protein and polynuclectide sequences of hopMilp-1 can be used in gene therapy and vaccine production. The hopMilp-1 community acids can be used as probes to detect, characterise and quantify compounded initial substrates for the recombinant engineering of hopMilp-1 compounded initial substrates for the recombinant engineering of hopMilp-1 compounded initial substrates for the recombinant engineering of hopMilp-1 compounded in a probes for the recombinant engineering of hopMilp-1 protein or proteins and for compounded in the proteins. The hopMilp-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hopMilp-1 proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as compounded in patients having specific deficiency in hopMilp-1 compounded and in vaccines or for replacement therapy. The concentration of disorder associated with the expression of hopMilp-1, in particular heart cand skeletal muscle disorders. hopMilp-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hopMilp-1 sequence data for this patent did not form part of the printed concentration, but was obtained in electronic format directly from WIPO at fifth the present invention. N.B. of the printed concentration, but was obtained in electronic format directly from WIPO at fifth the present sequence and obtained in electronic format directly from WIPO at fifth the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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Sequence 17 BP; 1 A; 4 C; 7

G; 5 T; 0 U; 0 Other;

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The present invention describes a human genome-derived myosin-like CC protein 1 (hcDMLP-1). The protein and polynucleotide sequences of hcDMLP-1 can be used in gene therapy and vaccine production. The hcDMLP-1 cC nucleic acids can be used as probes to detect, characterise and quantify comproved initial substrates for the recombinant engineering of hcDMLP-1 cC protein variants having desired phenotypic improvements, and for expressing the proteins. The hcDMLP-1 proteins or polypeptides may be cused as immunogens to raise antibodies that specifically recognise hcDMLP-1 proteins, as specific biomolecule and/or amount specifically of hcDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as characteris for surface-enhanced laser description ionisation, as concentration, and in vaccines or for replacement the concentration cC production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hcDMLP-1 may be used for diagnosing a cC disorder associated with the expression of hcDMLP-1, in particular heart cand skeletal muscle disorders. hcGMLP-1 is localised to chromosome 22. The present sequence in the expression of the present invention. N.B. The present sequence data for this patent did not form part of the printed cc specification, but was obtained in electronic format directly from WIPO at first wipo.int/pub/published_pct_sequence

8 Query Match Best Local & Matches Sequence 17 BP; 1 A; 1445 GGCCACCACACTGGGAG 1461 17 GCCAACCACACTGGGAG l Similarity 88.2 15; Conservative 0.5**%;** 88.2**%;** 5 C; 6 G; μ 0, Score 13.8; Pred. No. 8 5 T; Mismatches 88; DB 1; Length 17; <u>۲</u>. Indels 0 0

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Other;

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                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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; 2000US-0234687P.
; 2000US-0236359P.
; 2000GB-00024263.
; 2001WO-US000661.
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RESULT 176
ABNO8656
ID ABNO86
XX ABNO86
XX ABNO86
XX JP-MA1
XX Human
XX Human
KW muscle
XW skele
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Thes 15; Conserve
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                                              Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                  Human GDMLP-1
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Homo sapiens

RESULT 177
ABN00935
ID ABN009
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AC ABN009

ABN00935 standard; DNA; 17

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Cc The presents equence represents an oligomer used in the screening of the present sequence data for this patent did not form part of the printed capture wise, int/pub/mblished net sequence.
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30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
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                                                                                   0.5%;
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                                                                                     Score 13.8;
Pred. No. 88;
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ACAGCTGGAAGAGGCTG

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CC protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 cc nucleic acids can be used as probes to detect, characterise and quantify cc hGDMLP-1 nucleic acids in samples, as amplification substrates, to cc provide initial substrates for the recombinant engineering of hGDMLP-1 cc provide invariants having desired phenotypic improvements, and for cc expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP proteins, as standards in assays used to determine the concentration capture probes for surface-enhanced laser description ionisation, as cc therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The collar sequences encoding hGDMLP-1 may be used for diagnosing a cd disorder associated with the expression of hGDMLP-1, in paticular heart can disorder associated with the expression of hGDMLP-1, in paticular heart can disorder associated with the expression of hGDMLP-1, in paticular heart can disched a sequence of the expression of hGDMLP-1 is equence data for this patent did not form part of the printed content of the printed capture of the printed capture.
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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04-OCT-2000;
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or a
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as specific biomolecule capture probes for surface-enhanced laser
sorption ionization, comprises human myosin-like protein hGDMLP-1.
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Query Match Best Local S Matches 15

l Similarity

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88.2%;

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Length Indels

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RESULT 178
ABNO7863/c
ID ABNO7863;
XX 29-MAY-200
XX Human GDMI
XX Human; ger
KW muscle; my
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XX Human GDMI
XX 26-MAY-200
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The present invention describes a human genome-derived myosin-like CC protein 1 (hgDMLP-1). The protein and polynucleotide sequences of hgDMLP-1 can be used in gene therapy and vaccine production. The hgDMLP-1 culleic acids can be used as probes to detect, characterise and quantify checking in the protein probes to detect, characterise and quantify considering in the protein probes for the recombinant engineering of hgDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP cused as immunogens to raise antibodies that specifically recognise hgDMLP capture probes for surface-enhanced laser description ionisation, as capture probes for surface-enhanced laser description ionisation, as complement in patients having specific deficiency in hgDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production associated with the expression of hgDMLP-1, in particular heart can be seeded as the description in the particular heart contents and skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22.
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30-JAN-2001;
30-JAN-2001;
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27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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2000US-023458PP.

2000US-0235359P.

2001WO-US000661.

2001WO-US000662.

2001WO-US000664.

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RESULT 179
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ABQ634
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27-SEP-2000;
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23-MAY-2001;
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in KTC
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The invention relates to a novel isolated nucleic acid encoding human KTOM1 (kidney tumour overexpressed membrane) protein. The protein of the invention has cytostatic activity. The nucleotide may have a use in gene therapy. The KTOM1 nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1.
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n KTOM1 which
g., liver or
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ver or bone.
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2000US-0236359P.
2000GB-00024263.
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2001WO-US000662.
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rein, useful for treating subjects having defects
lfest as cancer of the kidney, or as a disorder of
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kidney; liver; bone marrow; brain; heart; lung;
al muscle; testis; uterus; placenta; probe; ss.
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New human kidney tumor overexpressed membrane (KTOM1) protein and nucl acids encoding the protein, useful for treating subjects having defect in KTOM1 which can manifest as cancer of the kidney, or as a disorder e.g., liver or bone.

n and nucleic ing defects disorder of

WPI; 2002-479509/51

therapy.

KTOM1 (kidney invention has

relates to a novel isolated nucleic acid ence tumour overexpressed membrane) protein. The prostatic activity. The nucleotide may have KTOM1 nucleic acids may be used to diagnose, t

protein e a use i

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or in human

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Example

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                                                                                                                                                           human kidney tumor overexpressed membrane (KTOM1) protein and nucles encoding the protein, useful for treating subjects having defections which can manifest as cancer of the kidney, or as a disorder
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ilarity 88.2%;
Conservative
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s to a novel isolated nucleic acid encoding human overexpressed membrane) protein. The protein of a atic activity. The nucleotide may have a use in g

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shared an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was

gimilar

SPA

Novel isolated human testis expressed Patched like protein (HTPL), use for identifying agonist and antagonist and specific binding partners.

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WPI; 2002-676582/73.

Example 2;

Page 110; 718pp; English.

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RESULT 182
ABV79112/c
ID ABV79912/c
ABV7912/c
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Matches 15
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-US000669.
09-OCT-2001; 2001US-0384761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        therapy. The KTOM1 nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1. Compositions comprising the nucleic acids, proteins or antibodies may used to treat subjects having defects in KTOM1 which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, by heart, lung, kidney, colon, skeletal muscle, testis, uterus and place function. The sequence represents a probe used in the invention to so the nt 1-1001 portion of human KTOM1a (ABQ63232)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mapped to human chromosome 10p12.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeuttc agents formale infertility and cancer. The present oligonucleotide was used in an example from the invention
                                           The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG), (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's Barcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; sturge Weber syndrome; leukaemia; ss; Osler-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
               vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, St
Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu
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                                                                                                                                                                            polynucleotide which down regulates expression of Ets-related gene, I for treating cancer, diabetic retinopathy, macular degeneration, itis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                              4; Page 127; 149pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ERG Amberzyme target sequence Seq ID No
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                                                                                                                   by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies curder conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or combined therapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically the regulate present nucleic acids, including antisense and enzymatic nucleic acid molecules which regulate expression of ERG, and cenzymatic nucleic acid molecules which regulate expression of ERG, and cenzymatic nucleic acid molecules which regulate expression of ERG, and
                                       Query Match
Best Local S
Matches 15
                                                                                                                 Sequence
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   1771
                                                           Similarity
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   AGGAGGAGGAGGCGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                    a patient having a condition associated with the
                                                                                                                   BP; 7
                                         Conservative
                                                                                                                 A; 1 C; 9
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                                                           Pred.
                                                                               Score 13.8;
                                                                                                                 0 T; 0 U;
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RESULT 184
ACC53896/c
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ACC53896;
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                  standard; DNA;
                                                             AGGAGGAAGAGGCAGAG
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tumour regression; apoptosis; virus cellular degeneration. tumour suppressor; antitumour; cytostatic; tumour suppression; suppressor sequence #2663 resistance; diagnosis

Human 27-JUN-2003

tumour

(first

entry)

FR2826373-A1 Homo sapiens

27-DEC-2002

20-JUN-2001; 2001FR-00008139

20-JUN-2001; 2001FR-00008139

(MOLE-) MOLECULAR ENGINES LAB SA.

Tuijnder M, Telerman Þ Amson

2003-250498/25

apoptosis eic acid sequences associated with tumor suppression, as or virus resistance are useful to diagnose and treat development of tumor cells and cell degeneration. regression, t viral

Claim ۲. Page 655; 798pp; French

with tumour suppression or regression, apoptosis or virus resinvention relates to these sequences or sequences having at leidentity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences invention is used to diagnose or treat viral disease or diseas characterized by development of tumour cells or cellular degen sequence represents an isolated nucleic acid sequence associated tumour suppression or regression, apoptosis or virus resistance. least The

osteoporosis

useful for

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CC The invention relates to a novel isolated 17 mer nucleic acid sequence, cc given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence with, after optimal cc alignment, at least 80 % identity to the 17 mer sequence that cc hybridizes to them under highly stringent conditions, or the complement co fany of them, or the corresponding RNA. The novel isolated nucleic acid so f the invention are useful as probes and primers for detecting, component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypoptides. Any of the nucleic acid, e.g. as one component of pharmaceutianing the nucleic acids, cells containing the vector or antibodies directed against the polypoptides are useful for preparation, specifically cancer but also Alpheimer's disease and compensation, specifically cancer but also Alpheimer's disease and compatient samples is useful for diagnosis and/or prognosis of these companies. The polypoptides can also be used to generate antibodies, and compatient samples is useful for diagnosis and/or prognosis of these companies. The nucleic acid sequences of the invention can be used in generate antibodies, and companies. The nucleic acid sequences of the invention can be used in generate antibodies, and companies of the sequences of the invention can be used in generate antibodies. The nucleic acid sequences of the invention can be used in generate components of protein companies. The polypoptide call sequences of the invention can be used in generate antibodies are useful as components of protein companies. The nucleic acid sequences of the invention can be used in generate antibodies and companies of the components of protein components.
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ABT34968
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 104; 720pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-2001; 2001FR-00011978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-2002; 2002WO-IB004208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABT34968;
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Pred. No. 8
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Sequence 17 BP; 5 A; 4 C;

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Matches 15
The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NPEB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg 2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-DEC-1992;
18-MAY-1994;
15-AUG-1994;
23-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cesphageal cancer; stomach cancer; colorectal cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; databetes; gemottabine, radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glionerulonephritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel enzymatic nucleic acid molecules which down regulates expression a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; Page 39; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-MAY-2001; 2001US-00864785.
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G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human;
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15; Conservative
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94US-00245466.
94US-00291932.
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Pred. No. 88;
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                                                                                                                                                                                                                                                                                                                                                                                                   New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
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developmental di
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proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are

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                                                 The present invention relates to novel human zinc finger-containing corporations and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is coded at chromosome 792.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome (MDZ7), and MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for dispnosing or monitoring a disease (CC caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The CC vaccines. The present sequence was used to illustrate the invention.
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chromosome 6p21.3-22.
developmental disorde
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7
6p21.3-22.2; chromosome 16p11.2; chromosome 15q
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encorner, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ61958;
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                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17
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Pred. No. 88
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ABZ66524,
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standard; RNA;

17

ВP

RESULT 191
ACD62072/c
ID ACD620
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XX 23-SEP
XX 23-SEP
XX W. Nuclei
KW RNA St
KW RNA St
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KW HBV re

23-SEP-2003 ACD62072

entry)

ACD62072 standard; RNA; 17

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HCV minus strand DNAzyme substrate sequence #383.

Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme amberzyme; G-cleaver ribozyme; decoy molecule; aptamer;

HBV reverse transcriptase; Enhancer I region; viral

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17

Matches Query Match Best Local

l Similarity

Conservative

0

0.5%; 88.2%;

Score 13.8; Pred. No. 88; Mismatches

DB 1; 2

Length

17

0

Gaps

0

1268

GGAAGAGGCTGAGGGCA 1284 GGAAGACGCTGAGGTCA

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acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ66216, ABZ664544 - ABZ65531, ABZ66520 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention
                                                                                                                                                                                                                                                                               Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encod HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
Sequence 17
                                                                                                                                                                                                                                                 Claim 4; Page 139; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ribozyme; short interfering RN; enzymatic nucleic acid; H-Ras; N-Ras;
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                                                                                                                                                                                                                  The invention relates to a novel short interfering RNA
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BP; 2 A; 7 C; 3 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        K-Ras;
                                                                                                                                                                                                    NA (siRNA) nucleic that modulates
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stabbility of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense cand more mucleic acid molecules include antisense cand more mucleic acids such as hammerhead ribozymes, DNAzymes, antisense are nucleic acid decoy molecules and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV by and or minus strand DNAzyme sequences disclosed in the present
                                                                             Query Match
Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2001; 2001US-00817879.
08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                         Sequence 17
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                                      882 ACCTTTGAGAGCCTGCA 898
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15; Conser
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MCSWIGGEN J.
MORRISSEY D.
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BLATT L.
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Roberts E;
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Pred. No. 8
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ACD65403 ID ACD XX

ACD65403 standard; RNA; 17

Query Match Best Local Similarity

0.5%; 88.2%; 5 C; 7

Sequence 17

BP; 5

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G; 0 T; 0 U; Score 13.8; Pred. No. 8

0 Other; DB 1;

Length 17;

RESULT 192

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or the synthesis, expression and/or stability of Hepatitis C virus (HCV) or CC Hepatitis B virus (HEV) RNA. The nucleic acid molecules include antisense and company of the cacids such as hammerhead ribozymes, DNAsymes, antisense cc inozymes, zinzymes, amberzymes, and G-cleever ribozymes, Also disclosed care nucleic acid decoy molecules and aptamers that bind to HBV reverse cc transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV CC compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and compounds of the invention are useful for the treatment of degenerative and clisease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carbon as cirrhosis, liver failure, and hepatocellular compounds or minus strand DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blatt L,
Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; ir amberzyme, G-cleaver ribozyme; decoy molecule; aptemer; amberzyme, G-cleaver ribozyme; tregion; viral repl
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08-UUN-2001; 2001US-00877478.
08-UUN-2001; 2001US-0296876P.
24-CCT-2001; 2001US-03350559.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                                                                                                                                                                                                                                        Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                             Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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MCSWIGGEN J.
MORRISSEY D.
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LEE P.
DRAPER K.
ROBERTS E.
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BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV;
                                                                                                                                                                                                                                                                                                                                                                        Page 311; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Macejak D,
Roberts E;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUN-2001;
08-JUN-2001;
24-OCT-2001;
The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amberzyme; G-cleaver ribozyme; decoy molecule; aptamer;
HBV reverse transcriptase; Enhancer I region; viral replication;
degenerative; disease state; HBV infection, HCV infection; cirrhosis;
liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zir
                                                                                                                                                                                                                                                                                                                       Claim 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                    Novel compound useful for hepatocellular carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-229207/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-)
(BLAT/)
(MACE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-OCT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   \vdash
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RIBOZYME PHARM
BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MACEJAK D.
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Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; 2001US-00817879.
; 2001US-00877478.
; 2001US-0296876P.
; 2001US-0335059P.
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                                                                                                                                                                                                                                                                                                                       267; 387pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                    treating cirrhosis, liver failure, or condition associated with hepat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis RNA stability; RNA expression; RNA synthesis, antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAyme; in amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; amberzyme; cleaver ribozyme; Iregion; viral replants reverse transcriptase; Enhancer I region; viral replants 
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08-UUN-2001; 2001US-00877478.
08-UUN-2001; 2001US-0296876P.
24-CCT-2001; 2001US-0335055P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                   Novel compound hepatocellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                       PAVCO P.
LEE P.
DRAPER K.
ROBERTS E.
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MORRISSEY
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                                                                                                                                                                                                                                                                                                                Macejak D,
Roberts E
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                                                                                                                                                         treating cirrhosis, liver failure, or condition associated with hepat
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Pred. No. 88;
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Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             anticonvulsant; antiparkinsonian; cerebroprotective; antimigraine;
antidepressant; analgesic; ophthalmological; vulnerary; antidiabetic;
dermatological; transgenic; chromosome 10q21.3; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADAMTS14; human; enzyme; neuroprotective; immunosuppressive; nootropic; antiinfertility; osteopathic; antiarthritic; antirheumatic; antiinflammatory, antiasthmatic; immunomodulator; antiallergic; cytostatic; antiulcer; vasotropic; antiarterioselerotic; cardiant;
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Hepatitis B virus (HBV
and enzymatic nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene exon 13 3' acceptor splice site.
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Pred. No. 8
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                                                                                 is an aminoprocollage processing. ADAMTS14
                                 present sequence is that of the 3' acceptor splice site of exon 13 of over human A Disintegrin and Metalloproteinase with Thrombospondin or repeats (ADAMTS) gene, denoted ADAMTS14, on chromosome 10q213. A A sequence for ADAMTS14 is given in ACC58643. ADAMTS14 (see ABR42736) an aminoprocollagen peptidase that functions in procollagen cessing. ADAMTS14 polynucleotides, polypeptides, vectors, cells nsfected by the vectors, and inhibitors directed against ADAMTS14 are d in the treatment and/or prevention of a range of diseases
                                                                                                                                                                                                                                  67pp; English.
                                                                                                                                                                                                                                                                                            s polynucleotide, useful fo of e.g. neurodegenerative,
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autoimmune, and
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밁 8 Query Match Best Local : Matches 1271 17 ACAGGCTGAGGCCAGAG 1 15; Similarity AGAGGCTGAGGGCAGAG Conservative 88.2%; 1287 0 Score 13.8; Pred. No. 88; Mismatches DB ۲. 2 Length Indels 0 Gaps 0

Sequence 17

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RESULT 196 ADB42204/c ID ADB42204 standard; 18-DEC-2003 04-DEC-2003 ADB42204; (revised) (first entry) DNA; 17

suppression/reversion associated nucleotide

diagnosis. virus resistance; cytostatic; primer; pro tic; antiviral; neuroprotective; nootropic; neuroleptic; ss; probe; tumour suppression; tumour reversion; apoptosis; esistance; transgenic animals; Alzheimer's disease; schizoph schizophrenia;

Homo sapiens.

WO2003040369-A2

15-MAY-2003

17-SEP-2002; 2002WO-IB004219

17-SEP-2001; 2001FR-00011981

(MOLE-) MOLECULAR ENGINES LAB

WPI; 2003-441574/41.

Telerman A,

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New useful e.g. polypeptide nucleic acid encoding human property for treatment of tumors and antibodies. prostate membrane-specific antigen, mors and viral infection, also relat related

Disclosure; Page 327; 771pp; French.

The invention relates to the isolation of 6327 nucleotide sequences, a fragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acides, as in vitro sense and antisense sequences, of nucleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce

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Analysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can also be used to screen for their specific interactive molecules, potentially useful for treating diseases associated with abnormal expression of the nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; inhibitor-kappa B-R; I-kappaBR; IXBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
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New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ი
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.8;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleotides'
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molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKRR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, vital or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; Page 75; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to antisense compounds targetted to
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Query Match
Best Local S
Matches 15
               93
                                 15;
4
                                          Similarity
               rescercerceasece 109
TEGECTCCTCCAGCACG
                                 Conservative
                                          0.5%;
88.2%;
20
                                 0
                                 Score 13.8; D
Pred. No. 1.3e
0; Mismatches
                                            .3e+02
                                                   DB 1;
                                                  Length 20;
                                   Indels
                                   0
                                   Gaps
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밁 S

Sequence

20

BP; 2 A; 9 C; 4 G; 5 T; 0 U; 0 Other;

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RESULT 198
ABLJ639
ID ABLJ63
XX
XX 22-APR
XX 22-APR
XX 1980sc
KW Human;
KW 1980sc
KW Hoddyki
KW novel
FW WO2001
XX
13-DEC
XX
XX
Claim
XX
XX
Claim
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Claim
CC The ir
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CC The intent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ss; lysosomal acid phosphatase 2; ACP2; gene; chromosome 11; lysosome-specific enzyme; orthophosphoric monoester hydrolysis; ledgkin's disease; HD; acid phosphatase deficiency; novel polymorphic site; ACP2 haplotype; ACP2 genotype; polymorphism; transgenic animal; primer; probe; primer-extension oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single nucleotide polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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SNP;

13-DEC-2001

07-JUN-2001; 2001WO-US018457

07-JUN-2000; 2000US-0210047P

(GENA-) GENAISSANCE PHARM INC.

WPI; 2002-154563/20

Kliem

SE,

Messer C,

Tanguay

Novel genetic variants of acid phosphatase 2, lysosomal pouseful in studying expression and function of the protein, screening drugs to treat diseases e.g. Hodgkin's disease. polypeptide in, and for gene

Page 14; 109pp; English.

The invention comprises the human lysosomal acid phosphatase 2 (ACP2) nucleic acid and protein sequences. Specifically, the invention relates to the discovery of 22 novel polymorphic siteally, the APC2 gene. The invention also comprises methods for haplotyping and genotyping the ACP2 gene in an individual. The ACP2 gene (located on chromosome 11) encodes a

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RESULT 199
ABL36360
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ss; lysosomal acid phosphatase 2; ACP2; gene; chromosome 11; lysosome-specific enzyme; orthophosphoric monester hydrolysis; lodgkin's disease; HD; acid phosphatase deficiency; novel polymorphic site; ACP2 haplotype; ACP2 genotype; polymorphism; transgenic animal; primer; probe; primer-extension oligonuclectide; SNP; single nucleotide polymorphism.
                                                         Novel genetic variants of acid phosphatase 2, lysosomal polypeptide gene useful in studying expression and function of the protein, and for screening drugs to treat diseases e.g. Hodgkin's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP;
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                                                                                                                                                                                                                                                                                                                       GENAISSANCE
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                                                                                                                                                                                                                                                 Messer C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                2000US-0210047P.
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14; 109pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 A; 6 C; 4 G; 2 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acid phosphatase 2 (ACP2) allele-specific PCR primer
                                                                                                                                                                                                                                                                                                                       PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92.9%;
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                                                                                                                                                                                                                                                        Tanguay DA
   English
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Ś B

The invention comprises the human lysosomal acid phosphatase 2 (ACP2) nucleic acid and protein sequences. Specifically, the invention relates to the discovery of 22 novel polymorphic sites within the APC2 gene. The invention also comprises methods for haplotyping and genotyping the ACP?

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ARTS487/C
ID AATS48
XX 25-MAR
AC AATS48
XX 25-MAR
DT 07-APR
XX Mouse
XX Enzyma
XX Enzyma
XX TNF-all
XX TNF-sel
XX TNF-sel
XX TNF-sel
XX TNS-sel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
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                                                                                                                                                                                                                23-FEB-1994;
29-MAR-1994;
04-APR-1994;
07-APR-1994;
                                                                                                                       15-APR-1994;
15-APR-1994;
18-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse relA hammerhead ribozyme target sequence (nt.
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07-APR-1997
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(first en
94US-00224483
94US-00227958
94US-00227958
94US-00245736.
94US-00271280.
94US-00291280.
94US-00291433
94US-00292620.
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94US-00218934.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13.6;
Pred. No. 70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 201
AXX569
ID AAX569
XX AXS69
AC AAX569
XX 16-OCT
DT 15-JUL
XX DB HIV-1
XX DNA-ta
KW DNA-ta
XX Viral
XX Human
                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                         á
                                                                                                                                                                                                                                                                                                                                                                The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves relA mRNA at the nucleotide base position indicated in the DE line. The relA gene product is a subunit of the transcriptional regulator NF-kappaB and is implicated specifically in the induction of inflammatory responses. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes are designed to cleave the target stepences and thereby inhibit relA expression, making them potentially useful for treating rheumatoid arthritis, restenosis and asthma as well as for increasing tolerance to transplanted trissues. The potential immunosuppressive properties of a ribozyme that cleaves relA mRNA means that uses are limited to local delivery, acute indications or ex vivo
                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-NIG-1994

08-SEP-1994

08-SEP-1994

23-SEP-1994

23-SEP-1994

28-SEP-1994

03-OCT-1994

11-OCT-1994

11-NOV-1994

10-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Grimm S,
Modak A,
Tracz D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ribozymes having modified bases and methods for producing them - in inhibiting disease related genes.
                          DNA-targeting conjugate; anticancer drug; viral DNA-cleaving viral DNA-binding agent; solid support; primer; ss.
                                                                       HIV-1 proviral
                                                                                                   16-OCT-2003
15-JUL-1999
                                                                                                                                             AAX56927;
                                                                                                                                                                          AAX56927
                                                                                                                                                                                                                                                                                                                                                                Sequence 15.BP; 2 A; 9 C; 2 G; 0 T; 2 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-351090/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-)
                                                                                                                                                                                                                                                                         1823 GGCCGGCGAGGTGGA 1837
                                                                                                                                                                                                                                              15
                                                                                                                                                                                                                                                                                                        14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 225; 407pp; English.
                                                                                                                                                                         standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Karpeisky A, Kisich K,
Pavco P, Beigleman L,
Usman N, Wincott FE,
                                                                                                                                                                                                                                                                                                                                                                                            (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dī,
                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                   (revised)
(first en
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94US-00321993.
94US-00334847.
94US-00337608.
94US-00345516.
94US-00357577.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94US-00311749.
94US-00314397.
94US-00316771.
                                                                       DNA fragment 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94US-00293520
94US-00300000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chowrira
                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                    93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ra B, Direnz
Kisich K,
                                                                                                                                                                                                                                                                                                        <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direnzo A, Draper KG, ch K, Matulic-Adamic J, n L, Sullivan SM, Sweed FE, Woolf T;
                                                                                                                                                                                                                                                                                                                      Score 13.4;
Pred. No. 76;
                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                 0 Other;
                                                                                                                                                                                                                                                                                                                                    DB 1; Length 15
                                                                                                                                                                                                                                                                                                        ۲.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sweedler D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           J. Dudycz IW;
J. Mcswiggen JA;
D. Thompson JD;
                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                         0
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RESULT 202
AA745448
ID AA7454
XX AA7454
XX AA7454
XX Bacter
XX Bacter
XX Bacter
XX Bacter
XX Bacter
XX Bacter
XX Argini
XW Argini
XX Bacter
XX Bacter
XX Argini
XX Bacter
XX Bacter
XX Argini

05-AUG-1997

(first

entry)

AAT45448

standard; RNA; 15

Bacteriophage lambda box B, RNA binding site.

Arginine rich; RNA; site; binding; box B; N protein; identification; screening; isolation; therapy; treatment; pathogenic; microorganism; disease; diagnosis; detection; ss.

멂 Ş

Query Match
Best Local Similarity
Matches 14; Conserv

Conservative

0

Mismatches

0

Gaps

0;

93.3%;

Score 13.4; Pred. No. 76;

멂

1; ۲.

Length Indels

1765 AAGATGAGGAGGAGG 1779

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CC phosphorodithioate, phosphoramidate or phosphoromidities group, OR1, SR1, CC an optionally substituted or modified nucleotide (N'), or an CC oligonucleotide of formula (N')gR2; g = 1-200; R1 = a protecting group; CC R2 = an H-phosphorame, alkanephosphonate, phosphortsetter, phosphite CC R2 = an H-phosphoramidate or phosphoramidate, phosphorodithioate, CC phosphoramidate or phosphoramidite group, OH, OR1, SR1 or CC phosphoramidate or phosphoramidite group, OH, OR1, SR1 or CC phosphorate, alkanephosphonate, phosphorothioate, phosphorate are compounds of formula CC groups and the group. Also mentioned are compounds of protecting group; and R4 = OH or an H-CC phosphorate, alkanephosphonate, phosphoratister, phosphoramidate group. Also claimed are new phosphoramidates, a grocess for preparing an oligonucleotide 5'-phosphate, a process for proparing an oligonucleotide 5'-phosphate, a process for preparing an oligonucleotide 5'-phosphate, a process for preparing an oligonucleotide 3'.5'-diphosphate. The CC proparing or -binding agents. The process for preparing or joingonucleotide of cleaving or -binding agents. The process for preparing oligonucleotide CC 1,5'-diphosphates is simple and suitable for use in automatic DNA CC genome, used to describe the method of the invention. (Updated on 16-OCT-CC 2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Derivatised solid supports and reagents for oligo:nucleotide synthesis and new oligo:nucleotide phosphoramidate conjugates.
Sequence 15 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -Z-CH2CH2-R, where: S' = a solid support; L = a bond or an (in)organic linker; Z = SO2 or S-S; R = OH, an H-phosphonate, alkanephosphonate, phosphorothioate, phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 44; 68pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Watanabe KA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9531434-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SLOK ) SLOAN KETTERING INST (ZWBI-) ZW BIOMEDICAL RES AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ren W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       describes novel derivatised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95WO-US006379
A; 0 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a solid
    G; 0 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CANCER RES
    0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         solid supports of formula S'-L
L = a bond or an (in)organic
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RESULT 203
AAZO70
XX
ID AAZO70
AC AAZO70
XX
O7-OCT
XX
DB Peptid
XX
CW Peptid
XX
W Peptid
XX
W Peptid
XX
W Peptid
XX
XX
Synthe
XX
SYnthe
XX
FH Key
FT modifi
                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                 Ś
                                                                                                                                                                                                                                                                                                                                                                        Bacteriophage lambda N protein residues 1-19, is an arginine rich RNA CD binding peptide (RBP) specific to the bacteriophage lambda box B, RNA binding site, i.e. the present sequence. It was used in a novel screening method, which comprised the construction of a vector encoding a hybrid protein, in which the 19 residue amino-terminal RBP of the phage lambda N protein was replaced by an arginine rich putative RBP. A second vector encoding the phage lambda termination site Nut and a marker gene was constructed, into which oligonucleotides containing box A of the Nut site and the appropriate RNA hairpin in place of box B were cloned. Plasmids were transformed into E. coli, and the expression of the reporter gene determined, showing that anti-termination was observed only with specific peptide/RNA interactions. The method can be used to isolate RBP, useful in therapy to block the life cycle of pathogenic microorganisms including viruses, e.g. HTV, and bacteria. The RBP can also be used to treat mammalian diseases resulting from impairment or loss of a natural RBP, as lead compounds for the development of the rapeutic compounds and in
                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Screening for RNA-binding polypeptide(s) - by expressing an antitermination protein fused to a test polypeptide and detecting exp of a reporter gene linked to a terminator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          stem_loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
                                                 Synthetic
                                                                        purification;
                                                                                                           Peptide
                                                                                                                                   07-OCT-1999
                                                                                                                                                                                  AAZ07075 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                         Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Fig 1A; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-012071/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (REGC ) UNIV CALIFORNIA
            modified_base
                                                                                                                                                                                                                                                                   1161 GCCCTGAAGAAGGCC 1175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ζ,
                                                                                  nucleic acid; PNA; polymer; solubility; modulation; synthesis;
                                                                                                           nucleic
                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                           GGCCUGAAGAAGGCC
                                                                                                                                                                                                                                                                                                                                                                e.g. pathogenic microorganism detection
                                                                                                                                                                                                                                                                                                                                         BP; 4 A; 4 C; 6'G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                  (first entry)
                                                                        analysis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lambda.
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/*tag=
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                                                                                                           acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SS,
                                                                                                                                                                                                                                                                                                      0.5%;
                                                                                                           oligomer #5.
                                                                                                                                                                                                                                            15
                                                                                                                                                                                                                                                                                                       Score 13.4;
Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Þ
                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                         76;
                                                                                                                                                                                                                                                                                                                   BG
                                                                                                                                                                                                                                                                                                                  <u>1</u>
                                                                                                                                                                                                                                                                                          1;
                                                                                                                                                                                                                                                                                                                  Length 15;
                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detecting expression
                                                                                                                                                                                                                                                                                          9
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Query Match
Best Local (
                                      Matches
                                                                                                                                               The present invention describes a branched composition (I) which is useful for improving the solubility of synthetic polymers (II) or all minimizing or eliminating self-aggregation of (II), where (II) is a nucleic acid (or analogue), peptide, peptide nucleic acid (PNA), polymenta or a linked polymer. Modification of (II) by (I) facilitate synthesis, purification and analysis of many insoluble polymers, and particularly purine-rich PNA polymers labeled with hydrophobic labels. The products can be used in research, diagnostic therapeutic applications. The present sequence represents a PNA used the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                     Branched compositions for improving the solubility of synthetic polymers or minimizing or eliminating polymer self-aggregation, particularly in peptide nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9937670-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                              Sequence 15
                                                                                                                                                                                                                                                                                                                                                                  Example 12; Page 40; 81pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gildea BD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-JAN-1998;
04-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BOST-) BOSTON PROBES INC
1767 GATGAGGAGGAGGAG 1781
                                      14;
                                                         Similarity
                                                                                                                BP;
                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0072772P
99US-00225048
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                                                                                                                5 A; 0 C; 10 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= b
/note= "g
uncharged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "g is modifie
carboxyfluorescein,
                                                       93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          λγειυοrescein, Ο
is an uncharged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         is modified to gether modifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          is modified
                                                                                                                                                                                                                                                                                                                                                                  English.
                                      0
                                                       Score 13.4;
Pred. No. 76;
                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to Flu-OE-g where Flu i
is 8-amino-3,6-dioxaoct
ether modifying moiety"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to g-E-NH2,
ying moiety"
                                                                         DB 1; Length
                                      1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             which
                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ,6-dioxaoctanoic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             is an
                                                                                                                                                                            diagnostic
a PNA used
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8
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ARESULT 204
AAF46416/c
ID AAF464
XX AAF464
XX AAF464
XX AATise:
DE IGFBP2
XX AATise:
KW Cytost.
KW Cytost.
KW Gytost.
KW IGF bi.
KW IGF bi.
KW GYOWth
KW GYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityviasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoes; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; keratosis; neoplasia; schooling the properties of the province of the pro
                                                                                                                                                                                                                                                                                                                                                                                                                                        Keratosis; neoplasia; scleroderma; wart; hyperneovascular condition; hyperplasia; neovascular condition of the retina; ss.
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(MURD-) MURDOCH CHILDRENS RES INST

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Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological, carddant, virucide; ophthalmological, keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

IGFBP2 oligonucleotide #23.

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Matches 14
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                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatclogical; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; lohthyosis; serborrhoea; ruba; keratosis; neoplasia; seleroderma; wart; skin cancer; sclerotic disease; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid tha inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15
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                                                                                                                                 21-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF45921 standard; DNA; 15
 WPI; 2001-041421/05
                                                                                                                                                                                                 WO200078341-A1
                                                                                                                                                                                                                                                              neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                               IGFBP2 oligonucleotide #760.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6; Page
                                                                  MURDOCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAAGCGGAGGAGGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           any other hyperplasia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                 2000WO-AU000693
                                 Werther
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                                                                                                 99US-0140345P
                                                                  CHILDRENS
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                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
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Pred. No. 76;
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                                    SR
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                                         The present invention relates to a method for ameliorating the effects of CC skin disorders. The method comprises contacting the skin with an CC antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense CC oligonucleotides of the present invention (see ARP45151 and ARF45153-CC oligonucleotides of the present invention (see ARP45151 and ARF45153-CC oligonucleotides of the present invention (see ARP45151 and ARF45153-CC oligonucleotides, ruba, pilaris, serborrhoea, keloids, keratosis, CC inthyposis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC neoplasias, scleroderma, warts, benign growths, cancers of the skin, a CC hyperneovascular condition such as a neovascular condition of the retina, CC disease, kidney disease, hyperproliferation of the inside of blood coversels or any other hyperplasia
Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ameliorating the effects of a disorder, UV (ultra-violet) treatment (optional) (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JV (ultra-violet) treatment (optional) and an antisense nucleinhibits or reduces growth factor mediated cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6; Page
  BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       39; 201pp; English
  4 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ф.
Q.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      psoriasis, by administering n antisense nucleic acid that ell proliferation and/or
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Matches 1530 CTGGAGGAGGCCAAG 1544 l Similarity
14; Conserv CTGGAGGAGCCCAAG 15 Conservative 0.5%; 93.3%; 0, Score 13.4; Mismatches ö. 76; B <u>-</u> ۲. Length Indels 0, Gaps

0

밁 S

30-MAR-2001 AAF45922; AAF45922 standard; (first DNA; 15 entry)

IGFBP2 oligonucleotide #761.

RESULT 208
AAF45922
ID AAF459
XX WO2000
XX WO2 Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

sapiens

WO200078341-A1

28-DEC-2000

21-JUN-2000; 2000WO-AU000693.

99US-0140345P

MURDOCH CHILDRENS RES INST

ر د Werther GA, Edmondson

Ameliorating the effects of a disorder, e.q UV (ultra-violet) treatment (optional) and inhibits or reduces growth factor mediated e.g. psoriasis, and an antisense ted cell prolifer by administering nucleic acid that ration and/or

inflammation

present invention relates to

method for ameliorating the effects

e F

6; Page 39;

201pp;

English

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RESULT 209
AAF45185
ID AAF451
XX AAF451
XX AAF451
XX AAF451
XX AAF451
XX AAF451
XX AF451
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XX A
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Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; opthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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                                                                                                                                                                           Ameliorating the effects
                                                                                                                                                                                                                                          WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUN-1999;
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                                                                                                JV (ultra-violet) treatment inhibits or reduces growth funhibits.
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                                                                                                                                                                                                                                                                                                  Werther
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                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0140345P
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                                                                                                                                                                                                                                                                                                  Edmondson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽P
                                                                                                                   a disorder, e.g. psoriasis, by administering (optional) and an antisense nucleic acid that factor mediated cell proliferation and/or
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Pred. No. 76;
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Example

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Page

34;

201pp; English

H-Ras, N-Ras,

inflammation.

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RESULT 210
ABZ665
XX
XX
ABZ665
XX
XX
ABZ665
XX
XX
ABZ665
XX
Z1-MAR
XX
Human;
XX
Human;
XX
Homo s
XX
C1-MAR
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C1-MAR
XX
CRIBO-TUN
PR 10-SEP
XX
PR 29-MAY
PR 29-MAY
PR 29-MAY
PR 10-SEP
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VIBO-TUN
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RESULT
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Best Local
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06-JUN-2001;
10-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 4 A; 2 C; 9 G; 0 T; 0 U; 0 Other;
The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Rass, H-Ras, N-Ras human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human HER2
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                                                                                                                                                                                                                                                                                                                                                                                                                Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                Claim 4; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic DNAzyme target #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001US-0294140P.
2001US-0296249P.
2001US-0318471P.
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                                                                                                                                                                                                                                                                                                                                                153; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA;
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Pred. No. 76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention

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RESULT 211
ACD82534
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Matches 14
          Query Match
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                                                                                The invention describes adaptor molecules, where each end of the adaptor is compatible with a nucleic acid digested with a restriction enzyme or a nucleic acid comprising an end that is compatible with a nucleic acid digested with a restriction enzyme. The adaptor molecules, compositions, kits and arrays are useful for cloning nucleic acid molecules that does not require the design and synthesis of oligonucleotides or PCR primers. The adaptors, kits and arrays are also useful for ligating two ends of a single nucleic acid molecule, or ligating two or more nucleic acid molecules. The kits can also be used for performing internal deletion molecules are ligated to a cloning vehicle, making the cloning procedure more rapid and efficient, and less error-prone. This sequence represents a nucleic acid cloning associated adaptor molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACD82534 standard;
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                                                                                                                                                                                                                                                                                           New adaptor molecules, useful for cloning nucleic acid molecules does not require the design and synthesis of oligonucleotides or
                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUN-2001; 2001US-00880313.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Adaptor molecule; nucleic acid cloning; nucleic acid ligating; internal deletion mutagenesis analysis; cloning vehicle; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic
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                                                                                                                                                                                                                                                                                                                                                          Flemington
                                                                                                                                                                                                                                                                                                                                                                                                            13-JUN-2001; 2001US-00880313.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-MAR-2003.
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                                                  Sequence 15
                                                                                                                                                                                                                                                     Claim 12; Fig 5; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                    (FLEM/) FLEMINGTON E K.
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            Similarity
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ilarity 93.3%;
Conservative
                                                 BP; 3 A;
 Conservative
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            93.3%;
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                                                  5 C; 5
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                                                  G; 2 T; 0 U; 0 Other;
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Pred. No. 76
               Score 13.4;
Pred. No. 7
             Pred.
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Mismatches
               76;
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                           DB 1;
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1247 GGTCCGGCTGCAGCA 1261

Homeo box consensus sequence primer (250)

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RESULT 212
ADA273
XX ADA273
XX ADA273
XX 20-NOV
XX 20-NOV
XX 10-NOV
XX LTAINSP
KW LTAINSP
KW LTAINSP
XX US2003
XX US2003
XX 11-NOV
XX INOK
PD 12-UNN
XX INOK
PD 12-UNN
XX INOK
PD 12-WN
XX INOK
PD 14-WN
XX INOK
P
RESULT 213
AAQ24927/c
ID AAQ249
XX
AC AAQ249
XX
DT 25-MAR
DT 19-NOV
XX
DE Homeo
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an oligonucleotide primer capable of specifically hybridising to a DNA having the sequence of the flanking regions of a microsatellite selected from M2-4-9, M2-2-12, M2-3-11, M2-2-20, M2-2-21, M2-2-22, M2-2-23, M2-2-24, M2-4-25, M2-4-26, M2-2-29, M2-2-32, M2-4-33, M2-4-37, M2-2-24, M2-4-26, M2-5-11, M2-2-46, and M2-2-48. The primer is useful for determining the number of repeat units of the microsatellite cited above. The primer is useful in HLA-related research, such as genetic mapping of HLA class II-associated diseases, transplantation matching, population genetics, and identification of recombination hot spots as well as linkage disequilibrium studies. The present sequence represents the human microsatellite repeat M2_3_3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ds; HLA-related research; HLA class II-associated disease; transplantation matching; recombination hot spot identific linkage disequilibrium study; human; microsatellite.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New oligonucleotide primer capable of specifically hybridizing to a DN having the sequence of the flanking regions of a microsatellite (e.g. M249), useful for HLA-related research, e.g. transplantation matching.
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                                                            25-MAR-2003
19-NOV-1992
                                                                                                                                                      AAQ24927;
                                                                                                                                                                                                                AAQ24927 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 5 A; 0 C; 10 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INOK/) INOKO H.
                                                                                                                                                                                                                                                                                                                                                                                                                            1767 GATGAGGAGGAGGAG 1781
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14;
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Similarity 93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                entry)
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Best Local Simi
Matches 14;
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                     17-APR-1991;
17-APR-1991;
17-APR-1991;
27-SEP-1991;
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29-JUL-1991;
                                                                                                                                                                                                                                                                                                                                Sequence 16 BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                       The selected primer is used in amplification reaction (SPAR).
                                                                                                                                                                                                                                                                                                                                                                               Claim 16; Page 39; 65pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-183683/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Cardineau GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single primer amplification; SPAR; ss
                                                                                                                                                                                                                                                                                                                                                                                              length data.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9207948-A1
                                                    08-APR-1991;
                                                                    23-NOV-1990;
                                                                                   25-NOV-1991;
                                                                                                 11-JUN-1992.
                                                                                                                  WO9209705-A1
                                                                                                                                 Synthetic
                                                                                                                                               Target; Human Immunodeficiency Virus; AIDS; triplex; hepatitis; herpes;
malignancy; ds.
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                                                                                                                                                                                       25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                             AAQ25457 standard; DNA; 16
                                                            18-JAN-1991;
      (GILE-)
                                                                                                                                                                                                                                                                               1877 GCAGCTGGAGGAGGA 1891
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                                                                                                                                                                                                                                                                                                          Similarity
      GILEAD
                                                                                                                                                                                                                                                                    GCAGCTGGAGAAGGA
                                                                                                                                                                                                                                                                                                                                                                                                     cid sequence single primer amplification - useful for genomic
analysis and polymorphism detection for restriction fragment
                                                                                                                                                                        HIV target duplex sequence.
                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                            Filner
                     90US-00617907.
91US-00643382.
91US-0068544.
91US-00686544.
91US-00686547.
91US-00686547.
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91US-00737919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91WO-US008233
      SCI INC
                                                                                   91WO-US008811.
                                                                                                                                                                                                                                                                                                          93.3%;
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                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                           Score 13.4;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                        (Updated
                                                                                                                                                                                                                                                                                                                                 σ
                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                            ,06
                                                                                                                                                                                                                                                                                                                                                          9 6
                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                          the single primer 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                   1.
                                                                                                                                                                                                                                                                                                                   Length 16
                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                    Gaps
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The sequence depicts a HIV viral duplex sequence which contains a purine-
rich region concentrated on one chain of the duplex. The sequence may be
prepd. by standard DNA synthesis. The HIV duplex sequence is used as a
target for novel oligomers which are capable of forming a triplex at
three such oligomers HIV141 -HIV143 are capable of forming a triplex with
this sequence. The oligomers are used in the diagnosis and therapy of HIV
infection. Similar oligomers may be used to target viral DNA duplexes
specific for hepatitis, herpes and malignancy. The triple helices form
under mild conditions thus assays may be carried out without subjecting
the test specimen to harsh conditions. The oligomer is able to inhibit
gene expression, as verified by in vitro systems See also AAQ25452-25501
and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 63; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                               New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Froehler B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krawczyk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matteucci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₹
                                                                                                                                                                                                                                                                                                                                                                                                                         which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Milligan
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                                        Query Match
Best Local
              1765 AAGATGAGGAGGAGG 1779
                                l Similarity
14; Conservat
2 AAGAGGAGGAGG 16
                                 Conservative
                                       0.5%;
93.3%;
                                 0
                                         Score 13.4;
Pred. No. 90;
                                 Mismatches
                                                  DB 1;
                                 1.
                                                  Length
                                  Indels
                                   0
                                   Gaps
                                   0
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Sequence 16 BP; 7 A; 0 C; 9 G; 0 T; 0 U; 0 Other;

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RESULT 215
AAQ70682/c
ID AAQ70682;
XX
AC AAQ70682;
XX
DT 25-MAR-2003 (revised)
DT 15-MAR-1995 (first entry)
XX
Exb-B2; upstream region; regu
XX
Exb-B2; upstream region; long
XX
Exb-B2; upstream region; regu
XX
Exb-B2; upstream regi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Erb-B2; upstream region; regulatory element; gene expression; triplex;
antisense; inhibition; screening; identification; cancer; breast cancer;
carcinoma; breast cancer; erythroleukaemia; sarcoma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Triplex forming oligonucleotide directed against Erb-B2 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
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94WO-US000348.

93US-00008897

Composition for decreasing gene transcription - co oligo:nucleotide or deriv. complementary to target get gene re region

Claim 12; Page 43; 71pp; English.

The Erb-B2 gene has a purine rich segment symmetry. This sequence, derived from the with substantial mirror Erb-B2 gene is located 69

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potential site of H-DNA formation. The overexpression of Erb-B2 is particularly associated with breast cancer. This triplex forming oligonucleotide directed against Erb-B2 and its derivatives may be used in the treatment of breast cancer, erythroleukaemia and sarcoma and more generally any disease involving the expression of Erb-B2. (Updated on 25-MAR-2003 to correct PN field.)

nucleotides upstream of the transcriptional start site and is

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RESULT 216
AAA95296
ID AAA952
XX AAA952
XX AAA952
XX Immuno
XW Immuno
XW inflam
XW conflu
XX WO2000
XX H-SEP
PN WO2000
XX 11-MAR
XX INHOF
XX INOIAT
PN INHOF
XX INOIAT
PN INOIAT
XX INOIAT
PT CRAM-2
PT CRAM-2
PT CRAM-2
YX Inolat
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Best Local S
Matches 14
                                                                                                        Matches
                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                 The present sequence is a PCR primer used during the identification of the murine confluency regulated adhesion molecule 2 (CRAM-2, also known as JAM-3) coding sequence. CRAM-2 is one of the vascular adhesion proteins of the immunoglobulin superfamily (Ig Sf). The CRAM-2 protein and coding sequence can be used in the treatment of cancer, inflammation, to doublate cell-cell interactions and angiogenesis, and in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 BP; 1 A; 10 C; 0 G; 5 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunoglobulin superfamily; Ig Sf; vascular adhesion molecule;
inflammation; cancer; wound; angiogenesis; mouse; JAM-3;
confluency regulated adhesion molecule 2; CRAM-2; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine CRAM-2 coding sequence identification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA95296 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-SEP-2000
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                                                                                                                                                                                                              Sequence 16
                                                                                                                                                                                                                                                                modulation of wound healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RMFD-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1777 AGGAGGCGGAGGAGG
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                                                                                                     1 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 15; 59pp; English.
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                                                  CACCCTCCGCACTCG 2257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   regulated
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                                                                                                                                                                                                                BP; 2 A; 9 C; 1 G; 4 T; 0 U;
                                                                                                     Conservative
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                                                                                                                               93.3%;
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Pred. No. 90;
                                                                                                                                 Score 13.4;
Pred. No. 9
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                                                                                                           Mismatches
                                                                                                                                    90;
                                                                                                                                                                                                                0 Other;
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                                                                                                                                                           DB 1;
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                                                                                                              Gaps
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RESULT 217
AAA953
XX AAA953
XX AAA953
XX IMMURINE
XX I
                                                                The present sequence is a PCR primer used to determine the amount of murine confluency regulated adhesion molecule 1 (CRAM-1, also known JAM-2) mRNA following assays differing in confluency. CRAM-1 is one the vascular adhesion proteins of the immunoglobulin superfamily (Ig The CRAM-1 protein and coding sequence can be used in the treatment cancer, inflammation, to modulate cell-cell interactions and angiogenesis, and in the modulation of wound healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunoglobulin superfamily; Ig Sf; vascular adhesion molecule; inflammation; cancer; wound; angiogenesis; mouse; JAM-2; confluency regulated adhesion molecule 1; CRAM-1; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Murine CRAM-1 mRNA transcript level
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Sequence 16
                                                                                                                                                                                                                                                                                                                                                                             Example; Page 16; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Imhof BA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aurrand-Lions M;
BP; 2 A; 9 C; 1 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DICTAGENE
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Query Match
Best Local S
Matches 14
 2243
                         Similarity
CACCCTCCCCACTCG 2257
                  Conservative
                         93.3%;
                  <u>.</u>
                          Score 13.4;
Pred. No. 9
                  Mismatches
                           90;
                                  DB 1;
                                 Length
                    Indels
                                    16;
                    0
                    Gaps
                    0
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RESULT 218
ABF360
ID ABF360
XX ABF360
XX ABF360
XX CAFEB
XX Oligon
XX SNP; G
XW Centra
XX Centra
XX Homo B
XX Homo B
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XX Homo B
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                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                Oligonucleotide SEQ ID NO 136087
                                                                                                                                                                         ABF36090;
                                                                                                                                             21-FEB-2002
                                                                                                                                                                                                    ABF36090 standard; DNA; 13
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                                                                                                                                                                                                                                                                             CACCCTCCTCACTCG
                                                                                                                                               (first
                                                                                                                                               entry)
                                                                                                                   for
                                                                                                                   detecting
                                                                                                                      SNP
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WO200177384-A2

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RESULT 219
ABF36091/c
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The cligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABT00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
           WPI; 2001-657177/75
                                     Olek A,
                                                                                        07-APR-2000; 2000DE-01019173
                                                                                                                    06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                         WO200177384-A2
                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                Oligonucleotide SEQ ID NO 136088 for
                                                                                                                                                                                                                                                                                                           21-FEB-2002
                                                                                                                                                                                                                                                                                                                                                               ABF36091 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 136087; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-657177/75
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                                                                EPIGENOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                                               GAGGAGGAGGAGG 13
                                     Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 0 C; 9 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                ĀG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%;
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                                        Berlin K;
                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 13
                                                                                                                                                                                                                                                                                   SNP TSC0033984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   typing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or peptide nucleic olymorphisms (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                    06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       methylation status.
                                                                       ABN83902
                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                          ABN83902 standard; DNA; 13
detection
                                                                                                                                                                                                                                     1770 GAGGAGGAGGAGG 1782
                                                                                                                                                                                                   13
                                                                                                                                                                                                                                                                      13;
                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                   GAGGAGGAGGAGG
                                                                                                                                                                                                                                                                                                                                                13 BP; 0 A; 9 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 136088; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                        0.5%;
ilarity 100.0%;
Conservative (
                                    (first
probe following target binding and extension.
                                  entry)
                                                                                                            ВP
                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                          Score 13;
Pred. No.
                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                           63 ;
                                                                                                                                                                                                                                                                                                           <u>ب</u>
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RESULT 220
ABN83902/c
ID ABN839
XX ABN839
XX ABN839
XX SNP; s
KW Sene e
XX Uniden
XX Uniden
XX EFT misc_f
FT misc_f

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                                                                                                           WPI; 2002-437468/47.
                                                                                                                                                                                                                                                                                                                             09-NOV-2000; 2000US-00710983
                                                                                                                                                                                                                                                                                                                                                                                                  08-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1207209-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single nucleotide polymorphism;
expression analysis; probe; ss.
                                                                                                                                                                                                                                                    AGILENT TECHNOLOGIES INC
                                                                                                                                                                                   DA,
                                                                                                                                                                                                                                                                                                                                                                                                     2001EP-00123992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /bound_moiety= "SNP containing target nucleic acit/note= "binds to bases 13-1 of the SNP containing nucleic acid (see ABN83901) following binding and extension of the probe"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    extension"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ρ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of probe prior to binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acid"
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Detecting a target using an addressable array of probes linked to substrate and observing the binding pattern is useful to determine

determine

single

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RESULT 221
ABN839
AC ABN839
XX ADN839
XX O6-SEE
XX SNP; s
KW SSP; s
KW Gene c
XX Unider
XX Unider
XX Unider
XX Unider
XX UNIDER
FT misc_1
XX Detect
PT nuclec
FT misclec
FT misclec
FT misclec
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to evaluating the presence of a target in an analyte, using an addressable array of probes linked to a substrate. The method comprises preparing a solution consisting of a buffer, target nucleic acid, DNA polymerase, deoxynucleotides and polynucleotide primers, then exposing the solution to the array so that the target hybridises, extending the bound probes, and observing a binding pattern on the array. The method is used to detect SNP's (single nucleotide polymorphisms) in diagnostic, screening, and gene expression analysis. The current sequence represents an SNP detection probe following target being a supplied of the surrent sequence represents an SNP detection probe following target being the surrent sequence represents an SNP detection probe following target
                                                                                                                                                                                                                                                                        22-MAY-2002
                                                                                                                                                                                                                                                                                                     EP1207209-A2
                                                                                                                                                                                                                                                                                                                                                         misc_feature
                                                         nucleotide
                                                                                                                                                                                                            09-NOV-2000; 2000US-00710983
                                                                                                                                                                                                                                         08-OCT-2001; 2001EP-00123992
                                                                                                                                                                                                                                                                                                                                                                                                                                        misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABN83901;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABN83901 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                    Amorese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single nucleotide polymorphism; diagnostic;
expression analysis; ss.
                                                                                                                    2002-437468/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13 CTCAGCTGTGCTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 Similarity
13; Conserv
                                                                                                                                                   DA,
                                                                                                                                                                                AGILENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCAGCTGTGCTC 1635
                                                         a target using an addressable array of probes linked to a and observing the binding pattern is useful to determine e polymorphisms in diagnostic, screening, and gene express:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative 0;
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                                                                                                                                                 Sampson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
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                                                                                                                                                                                TECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                           following
                                                                                                                                                                                                                                                                                                                                /note=
                                                                                                                                                                                                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                         /*tag= a
/bound_moiety= "SNP detection probe"
/note= "binds to bases 13-1 of the S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          representing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; 5
                                                                                                                                                                                                                                                                                                                                "location of SNP site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13
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                                                                                                                                                                                                                                                                                                                                                                           probe extension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 13; DB; Pred. No. 63; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnostic, screening, and gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP detection target.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 13; 63;
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                                                         and gene expression
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                                                                         to determine single
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                                                                                                                                                                                                                                                                                                                                                                                         probe
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Disclosure; Fig 3B;

17pp; English

Disclosure;

Fig

3E;

17pp;

English.

The invention relates to evaluating analyte, using an addressable array

the presence of a target in of probes linked to a substr

The

expression

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to evaluating the presence of a target in an analyte, using an addressable array of probes linked to a substrate. The method comprises preparing a solution consisting of a buffer, target nucleic acid, DNA polymerase, deoxynucleotides and polynucleotide primers, then exposing the solution to the array so that the target hybridises, extending the bound probes, and observing a binding pattern on the array. The method is used to detect SNP's (single nucleotide polymorphisms) in diagnostic, screening, and gene expression analysis. The current sequence represents a nucleotide fragment representing an SN detection target. The SNP located at position 7 of the fragment may be detected by a probe in the method of the invention
                                                                        Detecting a target using an addressable array of probes linked to a substrate and observing the binding pattern is useful to determine a nucleotide polymorphisms in diagnostic, screening, and gene expressions.
                                                                                                                              WPI;
                                                                                                                                                                                                                                 08-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                 misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13 BP; 1 A; 5 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                               (AGIL-) AGILENT TECHNOLOGIES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1623 CTCAGCTGTGCTC 1635
                                                                                                                            2002-437468/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single nucleotide polymorphism; diagnostic; screening; expression analysis; primer; ss.
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                                                                                                                                                       DA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTCAGCTGTGCTC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                    Sampson
                                                                                                                                                                                                                                 2001EP-00123992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer following extension of probe sequence
                                                                                                                                                                                                                                                                                                                                                               (see ABN83902)
primer"
                                                                                                                                                                                                                                                                                                    /*tag= a
/note= "original length of
probe and extension"
                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                          /bound_moiety= "SNP detection probe"
/note= "binds to bases 13-1 of the S
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                          to bases 13-1 of the SN ) following binding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                         primer following binding
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                                                                                                                                                                                                                                                                                                                                                                               SNP detection probe nd extension of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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AAV938 ID AAV938 AC AAV938
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-OCT-1997;
02-OCT-1997;
05-NOV-1997;
19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUN-1997;
03-JUL-1997;
22-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            extension of the probe sequence. Thi labels that may provide a means for present in the array
                                                                                                                                                                                                                                                                                                                                           Jarvis T,
Parry T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human B-raf
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                                                                                                     Claim
                                                                                                                                           Identifying new catalytic nucleic acid especially ribozymes that cleave Raf restenosis, and also new ribozymes and used as antiviral agents and synthons.
                                                                                                                                                                                                                                                                        WPI; 1999-009494/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-MAY-1997;
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                                                                                               179; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                      RIBOZYME PHARM INC
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                                                                                                                                                                                                                                                                                                               Beigelman L, M
                                                                                                                                                                                                                                                                                                                 Matulic-Adamic J, Reynolds M, Kisich K,
Beigelman L, Mcswiggen JA, Karpeisky A,
, Workman CT, Beaudry A, Sweedler D;
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97US-005171BP.
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97US-0051321P.
97US-0061324P.
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97US-0068212P.
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                                                                                                  English
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                                                                                                                                                                        that modulates selected processes RNA for treating cancer, modified nucleoside triphosphates
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A method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucle:

nucleic

muscular

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RESULT 22
AAX09460
XX AAX0
XX AAX0
XX AAX0
XX POly
XW Poly
XW dete
XX YOLY
XW GETE
XX HOMO
XX SYNT
OS SYNT
OS SYNT
OS HOMO
XX CLand
XX WPI;
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XX WPI;
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Best Local &
Matches 11
AAXO9121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X12937). These primers can be used in a method for determining polymorphic forms in an individual for use in e.9 forensics, paternity testing or for phenotypic typing for diseases such as agammaglobulinemia, disbetes insipidus, Lesch-Nyhan syndrome, musculi dystrophy, Wiskott-Aldrich syndrome, Fabry's disease, familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
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                                                                                                                                                                                                                                                                                                   New isolated nucleic acid segments from the human genome determining polymorphic forms for use in e.g. forensics, testing or phenotypic typing for disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phenotypic typing; characteristic; paternity testing; cherotypic typing; characteristic; infection; here disease; cancer; inflammation; drug; therapy; medimarker; primer; ss.
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                                                                                                                                                                                                                                               310pp; English.
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Pred. No. 77;
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; medicament;
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Best Local S
Matches 13
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Best Local Similarity
Matches 13; Conserv
                                                                                             The present invention relates to human immunodeficiency virus type 1 (HIV -1) O-type specific antigen and the polynucleotide sequence encoding it. The present sequence represents a PCR primer used in the methods of the
                                                                   Sequence 15
                                                                                                                                 Disclosure; Page 3;
                                                                                                                                                  HIV-1 O-type
                                                                                                                                                                                     Kim SY,
                                                                                                                                                                                                                                          14-APR-1998;
                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                primer;
                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus type 1; HIV-1 O-type specific antigen; PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15
                                                                                                                                                                    WPI; 2000-609488/58
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                                           Similarity
                                                                                       invention
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                                                                   BP; 4 A; 5 C; 6 G; 0 T; 0 U; 0 Other;
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                                 Conservative
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f) Mismatches
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Pred. No.
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92;
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hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willbbrand's disease, tuberous sclerosis, Ehlers-Danlos spherocytosis, beilders disease, tuberous sclerosis, Ehlers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria, autoimmune diseases, inflammation, cancer, diseases of the nervous system, infection by pathogenic microorganisms, and characteristics such as longevity, appearance (e.g. baldness, obesity), strength, speed, endurance, fertility, and susceptibility or receptivity to particular drugs or therapeutic treatments. The isolated polymorphic nucleic acid segments can also be used to produce medicaments for the treatment or prophylaxis of such diseases
AAS57269 standard; DNA; 15 BE XX AAS57269;

AC AAS57269;

AC AAS57269;

XX Human CHRNB2 allele specific XX Human; cholinergic receptor, KW CHRNB2, memory disorder; Alrk Chromosome 1921; schizophreni ADHD; autosomal dominant noct XX Homo sapiens.

XX W0200174833-A2.

XX HOMO sapiens.

XX W0200174833-A2.

XX 11-OCT-2001; 2001WO-US010666.

XX 03-APR-2000; 2000US-0217952P.

XX 03-APR-2000; 2000US-0217952P.

XX 03-APR-2000; 2000US-0217952P.

XX 03-APR-2000; 2000US-0217952P.

XX (GENA-) GENAISSANCE PHARM INC XX

Choi JY, Kliem SE, Koshy B, PR 13-JUL-2000; 2000US-0217952P.

XX (GENA-) GENAISSANCE PHARM INC XX

Choi JY, Kliem SE, Koshy B, PR 13-JUL-2000; 2000US-0217952P.

XX Genoryping cholinergic recept individual involves determini of nucleotide pair at polymory individual involves determini of nucleotide pair at polymory individual involves determini of nucleotide pair at polymory SX

Claim 15; Page 15; 82pp; Engl XX

Claim 15; Page 15; 82pp; Engl X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-APR-2001; 2001WO-US010666
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Genotyping cholinergic receptor, individual involves determining rgic receptor, nicotinic, beta-polypeptide 2 gene of an s determining for two copies of the gene, the identity at polymorphic sites selected from PS1-24. 03-APR-2000; 2000US-0194155P. 13-JUL-2000; 2000US-0217952P.

GENAISSANCE PHARM INC

Koshy B,

Lee HH,

Sanchis

P

Page 15; 82pp; English

The invention relates to genotyping/haplotyping the cholinergic receptor, CC micotinic, beta-polypeptide 2 (neuronal) (CHRNB2) gene of an individual, CC comprising determining for the two copies of the CHRNB2 gene present in CC the individual, the identity of the nucleotide pair at one or more CC polymorphic sites selected from PSI-24. Also include are oligonucleotides CC variants of CHRNB2. The method is useful for detecting novel CHRNB2 CC polymorphisms and for determining if an individual has a haplotype or CC polymorphisms and for determining if an individual has a haplotype or CC polymorphisms and for treating a specific condition or disease predicted to be associated with CHRNB2 activity (e.g. a memory disorder, Alzheimer's CC disease, epilepsy, a learning disorder, schizophrenia, attention CC defictlyhyperactivity disorder, Albab) and autosomal dominant nocturnal frontal lobe epilepsy (ADNFLE)), and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with CHRNB2 activity. The method is useful to screen for CC compounds targeting CHRNB2 to treat a specific conditions or disease predicted to be associated with CHRNB2 activity. The method is useful to screen for CC cassociated with CHRNB2 activity. The present sequence is an CC CHRNB2 gene is located on chromosome lq21. The present sequence is an CC chrnb2 gene is located on chromosome lq21. The present sequence is an method of the invention

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Š Query Match Best Local Matches 1778 Similarity GGAGGCGGAGGC 1792 Conservative 0.5%; 1, Score 13; DB Pred. No. 92; Mismatches DB 1; نر Length 15; Indels 0, Gaps 0;

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RESULT 226

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AAF483/C
ID AAF483
XX AAF483
XX 30-MAR
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                                                                                                                                                                                                                                                                                     oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; Inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
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RESULT 228 AAF53010/c ID AAF530

AAF53010

standard; DNA; 15

RESULT 229
AAF53012/c
ID AAF53012
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standard;

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                                                                                                                                                                                                                                                                                                  skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP), which is capable inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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[GF-I oligonucleotide #3972.

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RESULT 230
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                                                                                                                                                                                                                                                                                                               The present invention relates to a method for ameliorating the effects of cskin disorders. The method comprises contacting the skin with an cc antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 cc antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 cc inhibiting or reducing growth factor mediated cell proliferation, cc inflammation and/or other disorders. The present sequence is an coligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense coligonucleotides of the present invention (see AAF45151 and AAF45153-cc oligonucleotides of the present invention (see AAF45151 and AAF45153-cc ichthyosis, pityriasis, ruba, pilaris, serborrheea, keloids, keratosis, cc neoplasias, scleroderma, warts, benign growths, cancers of the skin, a complasias, scleroderma, warts, benign growths, cancers of the skin, a chyperneovascular condition of the retina, brain or skin, growth factor-mediated mallynancies, other sclerotic condisease, kidney disease, hyperproliferation of the inside of blood
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Homo

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

WO200078341-A1

sapiens

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28-DEC-2000

21-JUN-2000; 2000WO-AU000693

21-JUN-1999; 99US-0140345P

(MURD-) MURDOCH CHILDRENS RES INST.

GA,

Edmondson

WPI; 2001-041421/05

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or

Example 7; Page 55; 201pp; English.

The present invention relates to a method for ameliorating the effects of cskin disorders. The method comprises contacting the skin with an cantisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFs]-2 or IGFS93), which is capable of cinhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense coligonucleotide which can be used to design the antisense coligonucleotide which can be used to design the affects of psoriasis, CC oligonucleotides of the present invention (see AAP45151 and AAF48153-cc inchipyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC inchipyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC neoplasias, scleroderma, warts, benign growths, cancers of the skin, a cc brain or skin, growth factor-mediated malignancies, other sclerotic cc disease, kidney disease, hyperproliferation of the inside of blood cc vessels or any other hyperplasia

Sequence 15 BP; 0 A; 6 C; 6 G; 3 T; 0 ٦; 0 Other;

Query Match Best Local S Matches 13 l Similarity Conservative (0.5%; 0, Score 13; Pred. No. Pred. No. 92;
Mismatches (,, Length 15; Indels

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문 Ś 2121 15 CACGGGGCCGCAG 2133

RESULT 231 AAF52824; AAF52824

standard;

IGF-I oligonucleotide #3784.

entry)

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AC Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid, skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityrlasis; IGF binding protein; IGFBP-2; IGFBBB3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba. keloid;

antiproliferative; antiinflammatory; antipsoriatic;

disease;

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RESULT
AAF5282
Query Match
Best Local S
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             Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrheea; ruba; keratosis; meoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; hyperasorular condition of the retina; ss.
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skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBB]-2 or IGFBB3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide of the present invention (see AAF45151 and AAF45153-01igonucleotides of the spresent invention (see AAF45151 and AAF45153-01igonucleotides), ruba, pilaris, serborrhoea, keloids, keratosis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, aneoplasias, scleroderma, warts, bering growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
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effects

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Sequence 15 BP; 2 A; 4 C; 6 G; 3 T; 0 ς, 0 Other;

Query Match Best Local S Matches 13 Similarity Conservative 0.5%; 0 Score 13; Pred. No. ore 13; DB 1; Length 15; red. No. 92; Mismatches 0; Indels 0 Gaps 0

Ś 밁 1232 GCATGTGCTGGCA 1244 ω GCATGTGCTGGCA 15

RESULT 233
AAF53011/c
ID AAF530
XX AAF530
XX AAF530
XX AAF530
DT 30-MAR
XX IGF-I
XX ARTise
KW Cytost
KW Skin d
KW IGF bi
KW growth
KW keratc
KW hypern
KW hypern
KW hypern
KW homo s
XX
PN WO2000 30-MAR-2001 AAF53011; AAF53011 oligonucleotide #3971. standard; DNA; 15 (first entry) ₽P

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Antisense therapy, antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; opthbalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFB-3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; meoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

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Matches 13
                                                                                                                                                Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1, pityriasis; IGF binding protein; IGFBP-2; IGFBP3; Inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; hyperneovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see ARP45151 and ARP45153-G150). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kinney disease, hyperproliferation of the inside of blood
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21-JUN-2000; 2000WO-AU000693

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11-JAN-2001; 20-APR-2001; 24-APR-2001;

; 2001EP-00870005. ; 2001EP-00870085. ; 2001US-0286102P.

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09-JAN-2002; 18-JUL-2002

2002WO-EP000153

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ABZ34164/c
ID ABZ341
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AC ABZ341
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AC ABZ341
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AC HIV-1
AC HIV-1
XX
Human
KW Human
KW Probe;
XX Human
CS Human
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XX Human
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PF 09-JAN
XX 11-JAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a method for detecting mutations CC associated with anti-HIV drug resistance in a patient by detecting at C least one of the mutations K103N/R, V106A/I/L, V18EC/I, M18EV/I Y18EL, CC G199A/S/R, T215Y/F/D/S/A and/or Q151M/L in the reverse transcriptase (RT) CC of HIV strains in a biological sample using a specific set of probes CC optimised to function together in a reverse-hybridisation assay. The method and the nucleic acid sequences used in the method are useful for CC determining viral mutations and/or polymorphisms in the HIV RT gene associated with resistance. The probes are useful for CC detection, preferably in vitro detection of the mutations K103N/R, V106A/I/L, V138IC, Q15IM/L, M184VI, Y18BL, G199A/S/R and/or V106A/I/L, V138IC, I, Q15IM/L, M184VI, Y18BL, G199A/S/R and/or CC 1215Y/F/D/S/A in the RT of HIV strains in a biological sample, where the mutation is associated with anti-HIV drug resistance. The method provides a rapid, reliable and precise assay or determination and monitoring of CC antiviral drug resistance or mutations associated with drug resistance of viruses containing RT genes. ABZ33759 to ABZ34642 represent HIV RT sequences and probes which are used in the exemplification of the present of the present.
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Best Local S
Matches 13
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                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                     Aldehyde dehydrogenase 5 family member A1; ALDH5A1; succinate-semialdehyde dehydrogenase; gene therapy; primer; antisense technology; allele specific oligonucleotide; ASO; 4-hydroxybutyric aciduria; metabolic disease; transgenic an
                                                                                                                                                                                                                                                                                                                                                                                                  Aldehyde dehydrogenase 5 family, member
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13; Conserv
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                                                                                          Koshy B,
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                                                                                                                            PHARM INC.
                                                                                            Tanguay DA
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100.0%; Pred. No. 92;
ive 0; Mismatches
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New genetic variants of human aldehyde dehydrogenase 5 family, member Al ALDH5Al gene for treating metabolic diseases and for expressing ALDH5Al protein useful in identifying drugs to treat 4-hydroxybutyric aciduria.

Claim

15;

Page

13;

82pp;

English

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CC nuclectide sequence which is a polymorphic variant of a reference consequence for the aldehyde dehydrogenase 5 family, member A1 (succinate-semialdehyde dehydrogenase) (ALDH5A1) gene or its fragment. The comparising a sequence for the aldehyde dehydrogenase) (ALDH5A1) gene or its fragment. The comparison of the ALDH5A1 polymorphic variant with a candidate agent and assaying for binding activity. The polypeptide and haplotypes are useful for contentifying an association between a trait such as a clinical response to a drug targeting ALDH5A1 and a haplotype ALDH5A1 isogenes in vivo, are also useful for studying expression of the ALDH5A1 isogenes in vivo, for in vivo screening and testing of drugs against ALDH5A1 protein and compounds for 4-flydroxybutyric aciduria and metabolic diseases in a biological system. CC Antibodies are useful for immunoprecipitating the polypeptide from solution, for detecting ALDH5A1 protein isoforms in biological samples, immunohistochemical and immunofluorescence techniques. The polynucleotide is useful for gene therapy and antisense gene therapy. This sequence is colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphisms in the ALDH5A1 gene described in the method of the colymorphic colymorphic colymorphic colymorphisms in the ALDH5A1 gene described in the method of the colymorphic coly
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Matches 13
                                                                                                   New calmodulin-1 (CALM-1) isogene polymorphic variants, useful expressing CALM1 protein for use in screening for candidate drutreat diseases related to CALM1 activity such as Alzheimer's di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Calmodulin 1; CALM1; human; single nucleotide polymorphism; haplotyping; SCYA3; Alzheimer's disease; drug screening; calcium-dependent signal transduction; PCR primer; ss.
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CC The invention relates to an isolated polynucleotide comprising a sequence CC selected from a polymorphic variant of calmodulin 1 (CALMI). The CP colymorphic variant comprises an CALMI isogene defined by a haplotype CC selected from haplotypes 1-21 given in the specification. The polymorphisms are useful for studying the biological function of CALMI as CC well as in identifying drugs targeting this protein for the treatment of CC well as in identifying drugs targeting this protein for the treatment of CC adisorder related to its abnormal expression or function. The CC polymorphic variants may also be used in screening for compounds CC targeting CALMI to treat a specific condition or disease predicted to be associated with CALMI activity. Establishing CALMI haplotype or haplotype CC pair of an individual is useful for improving the efficiency and CC reliability of several steps in the discovery and development of drugs for treating diseases associated with SCYA3 activity, e.g. Alzheimer's CC disease and diseases involving defects in calcium-dependent signal CC transduction. Haplotyping the CALMI gene in an individual is also useful CC in the design of clinical trials of candidate drugs for treating a composition or disease predicted to be associated with CALMI activity. Asy55892-Asy59018 represent human CALMI allele-specific coligonucleotides and PCR primers of the invention
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24-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        probe;
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                                                                                                          mutations associated with anti-HIV drug resistance comprises at least one of the mutations in the HIV reverse transcriptase sing probes optimized to function together in a reverse-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                resistance;
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CC least one of the mutations K103N/R, V106A/I/L, Y181C/I, M184V/I, Y188L, CC G199A/S/R, T215Y/F/D/S/A and/or Q151M/L in the reverse transcriptase (RT) CC of HIV strains in a biological sample using a specific set of probes CC optimised to function together in a reverse-hybridisation assay. The method and the nucleic acid sequences used in the method are useful for CC determining viral mutations and/or polymorphisms in the HIV RT gene CC associated with resistance. The probes are useful for the genetic detection, preferably in vitro detection of the mutations K103N/R, CC V106A/I/L, Y181C/I, Q151M/L, M184V/I, Y188L, G190A/S/R and/or CC V105A/I/L, Y181C/I, Q151M/L, M184V/I, Y188L, G190A/S/R and/or CC T215Y/F/D/S/A in the RT of HIV strains in a biological sample, where the CC mutation is associated with anti-HIV drug resistance. The method provides CC a rapid, reliable and precise assay or determination and monitoring of CC viruses containing RT genes. ABZ33759 to ABZ34642 represent HIV RT conventions and probes which are used in the exemplification of the present of the present.
                                                                 invention
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Sequence 16 BP; 4 A; 1 C; 7
  G; 4 T; 0 U; 0 Other;
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Query Match
Best Local S
Matches 13
l Similarity
13; Conserv
0.5%;
ilarity 100.0%;
Conservative
٥,
          Score 13;
Pred. No.
 Mismatches
                  DB 1;
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                Length 16;
  Indels
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Gaps
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AAD44136;
                     AAD44136
                                                                                           1426 CCATCATCCACGT 1438
                                                                           16
                     standard; DNA; 15
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13-DEC-2002 primer #4 (first designed to bind human entry) MMP CATR region

Sequential consensus region-directed amplification; gene expression; disease diagnosis; gene analysis; human; matrix metalloproteinase; MMP; catalytic domain; CATR; PCR; primer; ss.

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ARBSULT 239
ARD44136
ID ARD441
XX ARD441
AC AR
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/note=
SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          base is given as
in the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          given as T in the sequence listing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence shown
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30-SEP-1998; 98US-00163485

03-OCT-1997; 03-OCT-1997; 97US-00943162. 97US-0108152P.

(-IVYU) UNIV VIRGINIA COMMONWEALTH INTELLECTUAL

Fillmore Ή Broaddus W, Gillies G;

WPI; 2002-412824/44.

Sequential consensus region-directed amplification for sorting mixture DNAs into 2 or more subsets or distinguishing gene expression patterns 2 samples, useful for disease diagnosis and gene analysis. inof

Example; Col 12; 19pp; English.

The invention relates to a method of sequential consensus region-directed amplification for sorting a mixture of DNAs into 2 or more subsets or

Claim 2 Page 26; 117pp; English

The present invention describes a method associated with anti-HIV drug resistance for detecting mutations in a patient by detecti detecting at

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RESULT 240
AAT90642/c
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recognition sequence; HCV; ribozyme; 5' untranslated region; nucleocapsid coding region; hairpin ribozyme; RNA cleavage; treatment; HCV infection; HCV contamination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      more subsets or distinguishing gene expression patterns in 2 samples e.g. for disease diagnosis and gene analysis. The present sequence is a PCR primer designed to bind to human matrix metalloproteinses (MMP) catalytic domain (CATR). This primer is used to illustrate the method of the
        AAT90621-650 represent recognition sequences found in the positive (-) strand of the Hepatitis C virus (HCV) RNA. The sequences are recognised by novel ribozymes which inhibit replication, infectivity or gene expression of HCV. The present sequence is located within the NS5 gene. Hairpin ribozymes of the present invention were designed based on
                                                                                                                                                                                                                                                                                                                                                                                     misc_feature
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                                                                             Example
                                                                                                                            Ribozyme(s)
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                                                                                                                  of viral
                                                                                                                                                                        Welch PJ,
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                                                                                                                 directed against hepatitis C virus f viral infection, and detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus
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Pred. No. 1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                 the CNR22 ribozyme"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for ribozyme cleavage
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                                                                                                                 - for prevention HCV contamination
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adjacent

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sequence recognition

The ribozymes

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ARXOSTAT 241
AAXOS6
AC AAXOS1
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CC spidrome (represented in AAX10269-X12937). These primers used in the commethod for determining polymorphic forms in an individual for use in e.g. forensics, paternity testing or for phenotypic typing for diseases such cas agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular cystrophy, Wiskott-Aldrich syndrome, Fabry's disease, familial cystrocytosis, von Willebrand's disease, tuberous sclerosis, hereditary cance, otherorynagic telangiectasia, familial colonic polyposis, Ehler-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria, appearance (e.g. baldness, obesity), strength, speed, can longevity, appearance (e.g. baldness, obesity), strength, speed, candurance, fertility, and susceptibility or receptivity to particular components can also be used to produce medicaments for the treatment or prophylaxis of such diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; Page 219;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic
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disease; cancer; inflammation;
marker; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ed nucleic acid segments from the human genome g polymorphic forms for use in e.g. forensics, phenotypic typing for disease.
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characteristic; infection; hereditary;
nflammation; drug; therapy; medicament;
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S Hepati
XX INFOCO
SX Hepati
XX 29-FEB
FR 27-FEB
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X
                                                                                                                                                        CC The present invention describes ribozymes (I) capable of inhibiting creplication, infectivity or gene expression of a hepatitis C virus (HCV), CC directed to target sequences AAA13438 to AAA13444, AAA13454 and AAA13465. CC (I) have virucide, hepatotropic and antiinflammatory activities. (I), or vectors comprising nucleotide sequences encoding (I), are useful for cells. (I) are useful for diagnosts, prevention and treatment of HCV cells. (I) are useful for preventing hepatitis C viral infection in a cells. (I) are useful for preventing hepatitis C viral infection in a cell. AAA13401 to AAA13405 represent examples of the briefest cell. AAA13401 to AAA13405 represent examples of the briefest comprisers used in the amplification of the HCV capsid sequence, AAA13408 to AAA13467 represent HCV hairpin ribozyme, AAA13408 and AAA13408 and AAA13407 represent holigonucleotides used in the construction of HCV hairpin ribozymes, all these sequences are used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SXX
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Best Local Simi
Matches 14;
                   Query Match
Best Local Similarity
  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ribozyme capable of inhibiting replication, expression of hepatitis C virus, useful for hepatitis C virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus; HCV; hairpin ribozyme; cleavage; recognition site; infection; virucide; hepatotropic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-FEB-1996;
27-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-MAR-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis C virus hairpin ribozyme recognition sequence SEQ ID NO:59
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                                                                                                  Sequence
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97WO-US003304
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Pred. No. 1.2e.
0; Mismatches
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                        Score 12.8; DB 1;
Pred. No. 1.2e+02
Mismatches
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No. 1
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                                                     is at risk of developing early-onset glaucoma comprising at least 1 (2) a kit for determining whether an individual is at risk of developing glaucoma with a high intraocular pressure at onset of disease, comprising at least 1 reagent that can be used to detect an ApoB4 allele in the individual; and/or at least 1 reagent that can be used to detect an ApoB4 allele in the individual, and/or at least 1 reagent that can be used to detect a T allele in the ApoB5 gene promoter. Identification of increased risk of glaucoma enables better treatment planning for affected individuals as well as for other family members who may be the affected individuals or disease gene carriers. The method provides a better and earlier means of diagnosis, so that preventative or palliative measures can be taken before significant damage to the optical nerve occurs
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TIGR) gene, or in TIGR gene promoter. The method comprises assessing the apolipoprotein E (ApoE) allele, or allele of the ApoE gene promoter. The invention also describes (1) a kit for determining whether an individual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for assessing the risk for developing early-onset glaucoma and for developing glaucoma with a intraocular pressure at the onset of disease in an individual havin mutation in a trabecular meshwork inducible glucocorticoid response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Assessing the risk of an individual for developing early-onset glaucoma, comprises assessing Apolipoprotein E alleles carrying a trabecular meshwork inducible glucocorticoid response gene mutation.
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Sequence 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             meshwork inducible glucocorticoid
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Query Match Best Local ! Matches

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Similarity

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Score 12.8; I Pred. No. 1.2e 0; Mismatches

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Length 16; Indels

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TCACCTTTGAGAGCCT

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ID AAF95038 standard; DNA; 16 BP
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AAF95038;
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DT 23-MAY-2001 (first entry)
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Mutant capture oligonucleotid
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Tubercle bacillus; drug sensi
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Tubercle bacillus; drug sensi
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C2-AUG-2000; 2000EP-00306563.
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O3-AUG-1999; 99JP-00220357.
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O3-AUG-1999; 99JP-00220357.
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CNISN) NISSHINBO IND INC.
AA (SYST-) SYSTEM RES INC.
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CNISN) NISSHINBO IND INC.
AA (SYST-) SYSTEM RES INC.
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CNISN) NISSHINBO IND INC.
AA (SYST-) SYSTEM RES INC.
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CNISN) NISSHINBO IND INC.
AA (SYST-) SYSTEM RES INC.
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RESULT 245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to oligonucleotides based on nucleotide sequences obtained from both wild-type tubercle bacilli (wtTB) that are consistent to a drug and mutant-type tubercle bacilli (mtTB) that are resistant to a drug. The drugs used in the present invention are creaistant to a drug. The drugs used in the present invention are creaistant to a drug. The drugs used in the present invention are creamburol (EB). The typeB gene is responsible for resistance to RFP; the crrs gene is responsible for resistance to SM and KM; the rpst gene is responsible for resistance to INH; the katG gene is responsible for resistance to INH; the katG gene is responsible for resistance to INH; cand the embB gene is responsible for resistance to EB. The present convention also relates to nucleic acid probes having part of a nucleotide sequence of tubercle bacilli (TB) responsible for drug resistance and convention can be used to enable the differentiation of drug resistance and the present invention can be used to enable the differentiation of drug resistance and the determination of infection with tubercle bacilli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         streptomycin; kanamycin; isoniazid; ethambutol; rpoB gene; rrs gene; rpsL gene; inhA gene; katG gene; embB gene; probe; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mutant capture oligonucleotide #31.
                                          16-JAN-2002
                                                                                                                                                                    AAS56903 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                    DNA;
                                          entry)
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RESULT 246
AAS15118/c
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5
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                                                                                                                                                                                                                                                                                                                                                                                                                sequences AAS56729-AAS56968 represent DNA encoding BRCA-1 regulators, ribozyme target recognition RNA sequences, DNA fragments encoding the RNP and primers used in the methods of the invention. Hybridisation of ribozymes to their targets results in cleavage of the RNA target. The ribozymes can be used to cleave regulators of the tumour suppressor BRCA-1, resulting in upregulation or downregulation of BRCA-1 in a cell. The mRNA targets include those encoding the BRCA-1 regulator BRC1 inhibitor dominant negative 4 (ID4), breast basic conserved protein 1 (BBC1), GHLR2, AF6, BR2 and BR3. Regulation of BRCA-1 is useful for treating and diagnosing cancer and other proliferative disorders. The severity of an incidence of cancer can be lessened by regulating tumour proliferation through modulation of BRCA-1 expression. The sequences of the invention are useful in the development of anti-cancer drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; BRCA-1 regulator; ribozyme; BR1; RNA target recognition; probe; cytostatic; RNA cleavage; tumour suppressor; PCR primer; CHLR2; AF6; B inhibitor dominant negative 4; breast basic conserved protein 1; BBC1; BR3; ID4; cancer; proliferative disorder; tumour proliferation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Validation ribozyme DNA sequence
                                   Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16 BP; 1 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                               Human; ss; CYI hybridisation
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                                                                                                                                                    16-JAN-2002
                                                                                                                                                                               AAS15118;
                                                                                                                                                                                                          AAS15118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel polypeptides that are the regulators of BRCA-1, useful for treating cancer and diagnosing the presence of neoplastic cells in biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-611503/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Beger C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (IMMU-)
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                                                                                                                       HyBeacon
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BEGER C.
                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                          standard; DNA; 16
                                                                                                                                                                                                                                                                                CCTGCGCAGACGGCAG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Barber J,
                                                                               CYP2D6; cytochrome P450; SNP; single ion beacon; 2D64E; F HyBeacon probe; I
                                                                                                                       probe
                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                    (first entry)
                                                                                                                         for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97pp; English.
                                                                                                                                                                                                                                                                                                                                                 0.5%;
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                                                                                                                         human CYP2D6,
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                          2D64E
                                                                                                                                                                                                                                                                                                                                                    .2e+02
                                                                                                                                                                                                                                                                                                                                                                 DB 1;
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                                                                                  nucleotide polymorphism;
pna-RNA hybrid.
                                                                                                                                                                                                                                                                                                                                                              Length 16;
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Key modified_base

Location/Qualifiers

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RESULT 247
AASI5115/c
ID AASI51
AC AASI51
XX ASI51
XX I6-JAN
XX Human;
XX Human;
XX Homo &
OS Synthe
XX Synthe
XX Hodifi
FH Key
FH Key
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Best Local
                                                                                                                                                                                                                                                                                                                                                                               oligonuclectide having substantially no secondary structure, and formed of nuclectides, one of which is labeled with a reporter, and no associated quencher. The beacon is used to detect, identify or quantify a target sequence in a sample, and to differentiate between homozygous and heterozygous polynuclectide targets. The present sequence is an F-Q HyBeacon probe targetting the a human gene for cytochrome P450, CYP2D6 which is known to contain several single nuclectide polymorphisms (SNP) and is used to demonstrate the use of the hybridisation beacons of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A hybridization beacon which is a single stranded oligonucleotide labeled with a fluorophore is useful to discriminate between polymorphic variants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAR-2000;
02-NOV-2000;
                                                                                                                                                                                                                                                                                                                                             Sequence 16 BP; 0 A; 3 C; 10 G; 2 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-616532/71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          French
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                                                                        Human; ss; CYP hybridisation
                                                                                                           F HyBeacon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 27; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (LGCT-)
Key
modified_base
                                     Homo sapiens
Synthetic.
                                                                                                                                     16-JAN-2002
                                                                                                                                                             AAS15115;
                                                                                                                                                                                   AAS15115 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                     invention in detecting the SNPs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  target
                                                                                                                                                                                                                                                                     1704 CAGCCCCAGGAGGCCC 1719
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                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotides.
                                                                        CYP2D6; cytochrome P450; SNP; single nucleotide polymorphism;
ion beacon; 2D64C*; F HyBeacon probe; DNA-RNA hybrid.
                                                                                                           probe
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                                                                                                                                                                                                                                                                                                Conservative
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2000GB-00026749
                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   relates to a hybridisation beacon which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= p
/mod_base= OTHER
/~~te= "U is cova
  Location/Qualifiers
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16
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/mod_base= OTHER
/note= "The 3' end of the probe is blocked with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     octanediol group"
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                                                                                                            for human
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87.5%;
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                                                                                                            CYP2D6,
                                                                                                                                                                                    ВP
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                                                                                                                                                                                                                                                                                                           Score 12.8; DB 1;
Pred. No. 1.2e+02;
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                                                                                                              2D64C*.
                                                                                                                                                                                                                                                                                                                      Length 16;
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RESULT 248
ABK4064
ID ABK406
XX ABK406
XX 21-MAY
XX 21-MAY
XX 21-MAY
XX Cardia
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                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                           oligonucleotide having substantially no secondary structure, and formed of nucleotides, one of which is labeled with a reporter, and no associated quencher. The beacon is used to detect, identify or quantify a target sequence in a sample, and to differentiate between homozygous and heterozygous polynucleotide targets. The present sequence is an P-Q HyBeacon probe targetting the a human gene for cytochrome P450, CYP2D6 which is known to contain several single nucleotide polynucleotide CNP) and is used to demonstrate the use of the hybridisation beacons of the invention in detecting the SNPs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A hybridization beacon which is with a fluorophore is useful to of target oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                         Sequence 16 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 27; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        French DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-MAR-2000;
02-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (LGCT-) LGC TEDDINGTON LTD.
                                                                                                                                                                        ABK40654 standard; DNA; 16
                                                                                                                                                                                                                                                      1704
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                                                                                                                                                                                                                                                                              14;
                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                      CAGCCCCAGGAGGCCC 1719
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                                                                                                                                                                                                                                                                             Conservative
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2000GB-00026749
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/label=
16
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/mod_base= OTHER
.... : COV
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/note= "The 3' e
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                                                                                                                                                                                                                                                                                                                         A; 3 C; 10 G;
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                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                              Score 12.8; DB 1;
Pred. No. 1.2e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       covalently linked to a FAM fluorophore"
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f
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                                                                                                                                                                                                                                                                                                                            Ç;
                                                                                                                                                                                                                                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                2
                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                       16;
                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                  Gaps
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variants
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ss; antisense; betal adrenoceptor; betal-AR; vasotropic; cardiant; hypertension; hypertrophy; cardiac ischaemia; cardiovascular disease; cardiac dysfunction.

WO200204623-A2

Human betal-adrenoceptor antisense

oligonucleotide #76.

hypotensive;

0

21-MAY-2002 ABK40654;

(first entry)

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RESULT 249
ABS98338/c
ID ABS98338/c
ID ABS983
XX
AC ABS983
XX
DE Human
XX
Human
XX
Human
XX
KW Cytoch
KW aryl h
KW aryl h
KW cytoch
KW cytoch
KW aryl b
KW cytoch
KW cytoch
KW aryl b
KW cytoch
KW aryl b
KW cytoch
KW Gylicat
KW Gylicat
KW HNMT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to an isolated antisense oligonucleotide of 9-35 cc nucleotides in length, which specifically binds to a portion of an mRNA ceppressed from a gene encoding a mammalian betal-adrenceptor (AR) cpolypeptide and alters the translation of the mRNA nito the betal-AR cc polypeptide in a host cell expressing the mRNA. Also included are a crecombinant vector comprising the antisense oligonucleotide, and a host cell comprising the vector. A composition comprising the antisense oligonucleotide, and a host cell comprising the vector. A composition comprising the antisense in treating or ameliorating hypertension, hypertrophy and cardiac ischaemia in a mammal. A composition comprising the antisense oligonucleotide is also useful for reducing the level of betal-AR polypeptide, the cell, particularly human cell, and for decreasing blood pressure in a cmammal, where the antisense oligonucleotide is operably linked to a promoter capable of expressing the oligonucleotide in the cell. A composition comprising a selected nuclei acid segment that comprises a full-length, or is a full length betal-adrenceptor antisense collymucleotide in a cell is also useful for reducing the level of betal-AR expressing the oligonucleotide in the cell. A composition in a cell is also useful for reducing the level of betal-AR adrenceptor polymetide in a mammalian host cell. The antisense colligonucleotide is also useful for other cardiovascular diseases and cardiac dysfunction in humans. The present sequence is a betal-AR mRNA targetting antisense oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local (
Human, ss; primer; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1; PCR; cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF; adrenergic receptor beta1; ADRR1; aryl hydrocarbon; AHR; MRP3; NR112; aryl hydrocarbon receptor nuclear translocator; ARN; cathepsin S; CTSS; cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological; epoxide hydroxylase 2; EPEX2; 5-lipoxygenase activating protein; FLAP; glutathione-S-transferase; 12; GST12; histamine-N-methyl transferase; NMMT; Kallikrein 2; KIK2; nicotinamide-N-methyl transferase; NMMT; NADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense oligonucleotides that specifically bind to mRNA encoding beta 1-adrenoreceptor polypeptide, useful for treating cardiac diseases of the cardiovascular diseases of the cardiovascular diseases of the cardiovascular diseases of the cardiovascular diseases
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                                                                                                                                                                                                                                 Human multidrug resistance associated protein 3 PCR primer
                                                                                                                                                                                                                                                                              23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                       ABS98338 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 2 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                     DNA; 16
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Pred. No. 1
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UDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7; UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uP multidrug resistance 1; lactotransferrin; crphan nuclear receptor; multidrug resistance associated protein 3; cancer; prostate; acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5; altered drug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological.
28-NOV-2001; 2001WO-US044838.
                                                                                                                                                                                   WO200257410-A2
                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               uPA;
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disorder-related traits Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for

Guida M,

Hall SCI LAB

2002-698522/75

(DNAS-)

28-NOV-2000; 2000US-00724389.

Example 24; Page 150; 714pp; English

comblecule comprising at least one base variation from that of a known comblecule comprising at least one base variation from that of a known controlled comprising at least one base variation from that of a known controlled comprising at least one base variation from that of a known controlled comprising at least one base variation from that of a known controlled comprising the problem controlled comprising the problem (PARP) (CTSS), cyclooxenase 2 (COX2), diazepam binding conhibitor (DBI), epoxide hydroxylase 2 (EBHX2), 5-lipoxygenase activating conhibitor (DBI), epoxide hydroxylase 2 (COX2), diazepam binding contransferase (HMMT), KADIH quinone oxidoreductase 2 (MOC2), histamine-N-methyl contransferase (HMMT), MDP-glucuronosyl transferase 2B4 (CC (UGT2B15), urckinase receptor (URA), multidrug resistance associated protein 3 (CMRCH2), UDP-glucuronosyl transferase 2B4 (CC (UGT2B15), urckinase receptor (URA), multidrug resistance i (CMRCH2), UDP-glucuronosyl transferase 2B4 (CC (UGT2B15), urckinase receptor (URA), multidrug resistance i (CMRCH2), UDP-glucuronosyl transferase 2B4 (CC (UGT2B1), urckinase receptor (URA), multidrug resistance i (CMRCH2), UDP-glucuronosyl transferase 2B4 (CC (URA), urckinase receptor (URA), multidrug resistance i (CMRCH2), URA), urckinase receptor (URA), urckinase receptor (URA) This invention relates to the sequence of an isolated nucleic acid

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T; 0 U;
0 Other;
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Best Local Similarity Query Match 87.5**%**; Score Pred. 12.8; No. 1 DB 1; .2e+02; Length 16;

Human

GDMLP-1 17-mer scanning

SEQ ID

NO:5 sequence

SEQ

ID NO:8648.

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ABN08656/
ID ABN0
XX
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DT 29-M
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ABX74378/c
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                                                                                RESULT 251
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Best Local S
Matches 14
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                                                                                                                                                                                                                               The invention relates to a new ribozyme with the ability to inhibit replication, infectivity or gene expression of a Hepatitis C Virus (HCV) by cleaving the positive strand genomic RNA of HCV at a sequence having 16 bp. Also included are a nucleic acid encoding the ribozyme, a host cell containing the ribozyme or vector, a vector comprising a promoter operably linked to the nucleic acid, producing a ribozyme, interfering with HCV replication or gene expression in a cell infected in a cell culture with HCV or a composition comprising the ribozyme and a carrier or diluent. The ribozyme is useful for treating or preventing HCV infection. The present sequence is an HCV -ve strand recognition sequence for a ribozyme of the invention
                                                                                                                                                                                                                                                                                                                                                                                                New ribozyme having the ability to inhibit replication, infectivity or gene expression of a Hepatitis C Virus (HCV), useful for treating or preventing HCV infection.
                                                                                                                                                                                                          Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hairpin ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C recognition sequence
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             29-MAY-2002
                                    ABN08656;
                                                                                                                                                                                                                                                                                                                                                                           Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barber JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-FEB-1996;
20-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus
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                                                         ABN08656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (IMMU-) IMMUSOL INC.
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                                                                                                                                        880 TCACCTTTGAGAGCCT 895
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                                                          standard;
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                                                                                                                                                                                                           BP; 5 A; 2 C; 5 G; 0 T; 4 U; 0 Other;
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97US-00954210
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                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                         0.5%;
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Pred. No. 1.2e
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1652

CCAGCTGCAGAGGCAG 1667

Query Match
Best Local Similarity
Matches 14; Conserv

Conservative

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Score 12.8; DB 1; Pred. No. 1.4e+02; 0; Mismatches 2;

<u>ب</u>

0

Gaps

0

Length 17; Indels

87.5%;

17

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A; 5 C; 6

G; 3

7; 0

4

0 Other

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The present invention describes a human genome-derived myosin-like protein 1 (hgDMLP-1). The protein and polynucleotide sequences of hgDMLP-1 can be used in gene therapy and vaccine production. The hgDMLP-1 curvate is used as probes to detect, characterise and quantify comproved initial substrates for the recombinant engineering of hgDMLP-1 provide initial substrates for the recombinant engineering of hgDMLP-1 comprovements, and for proved in variants having desired phenotypic improvements, and for expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hgDMLP proteins, as specific biomolecule capture probes for surface-enhance laser described expecting and in vaccines or for replacement therapy. The polymucleotide sequences encoding hgDMLP-1 may be used for diagnosing a consolited with the expression of hgDMLP-1 in particular heart and skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22 or he present sequence represents an oligomer used in the screening of the present invention. N.B.

The sequence data for this patent did not form part of the printed contribution in the semiplification of hgbmLP-1 wino, int/onl/hubleided not sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                 New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
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                           .wipo.int/pub/published_pct_sequence
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2000US-0234687P
2000US-0236359P
2000US-02024263
2001WO-US000661
2001WO-US000663
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2001WO-US000665.
2001WO-US000666.
2001WO-US000667.
2001WO-US000668.
                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID
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2001WO-US000670.
2001US-0266860P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INC.
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                                                                                                                                                                                                                                                                                                                                                                                                    214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rank DR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shannon
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CCAGCTGCAGCTGCAG 1

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AAL61563
                         The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-NOV-2002; 2002WO-US035597.
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ikappab r; antisense; immune response; infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-SEP-2003
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                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-NOV-2001; 2001US-00993731.
                                                                                                                                                                                                                                                                          Claim 3; Page 74; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003042360-A2
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tumour; prophylaxis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                         2003-468635/44.
                                                                                                                                                                                                                                                                                                       response or
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16. .2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note= "2'-methoxyethyl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (2'-MOE) nucleotides
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Seguence 20

BP; 4 A; 6 C;

7 G;

3 T;

0 U; 0 Other

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1227 CTCCAGCATGTGCTGG 1242

Conservative

0,

0.5%;

Score 12.8; Pred. No. 1 Mismatches

1.9e+02 멂

Length Indels

20

0,

Gaps

0

Query Match
Best Local Similarity
Matches 14; Conserv

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RESULT 253
ABK9982/C
ID ABK998
XX ABK998
XX 1-0C1
XX 21-0C1
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                                                 The invention describes a compound (I) 8-50 nucleobases in length cargeted to a nucleic acid molecule (II) encoding RAIDD which is an CC targeted to a nucleic acid molecule (II) encoding RAIDD which is an CC adaptor molecule containing both death domain (DD) and caspase crecitiment domains (CARD), where (I) specifically hybridises with and CC inhibits expression of RAIDD, or specifically hybridises with at least an CC inhibiting the expression of RAIDD (Receptor interacting protein (RIP) associated ICH-1/CED-3-homologous protein with death domain) in cells or CC tissues, and for treating an animal having a disease or condition CC associated with RAIDD, where the disease or condition is a hyperproliferative disorder such as cancer, or a growth or metabolic disorder. (I) is also useful for diagnostics, therapeutics, prophylaxis, as research reagents and kits, for distinguishing functions of various anembers of a biological pathway, and in antisense gene therapy. (I) is calso useful prophylactically, e.g. to prevent or delay infection, calso useful prophylactically, e.g. to prevent or delay infection. RAIDD antisense oligonucleotide used to control expression of the RAIDD protein
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CARD; hyperproliferative disorder; cancer; growth disorder; mouse; metabolic disorder; infection; inflammation; tumour formation; RIP associated ICH-1/CED-3-homologous protein with death domain; receptor interacting protein; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel antisense compound that hybridizes and inhibits nucleic acid encoding RAIDD which is an adaptor molecule containing both death domain and caspase recruitment domains, for treating hyperproliferative
Sequence
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappaB-B-related, ikappab r, nuclear factor of kappa Iijht polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
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                                                                                                                                                                                                                                                                                New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor kappa B-R (also known as I kappaBR, IKBR, I-kappa-B-related, ikappabr, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parastito.) They are useful for diagnostics, therapeutics, viral or parastic.) They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour
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                                                                           New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a height immune response or infection.
The invention relates to antisense compounds targetted to a nucleic admolecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappaB r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to
                                                                                                                                                                                                                            05-NOV-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human inhibitor-kappa B-R antisense oligonucleotide,
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                                                         Claim
                                                                                                                                                                                                       13-NOV-2001;
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                                                                                                                                    2003-468635/44
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                                                                                                                                                           Watt
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                                                                                                                                                                                  PHARM
                                                                                                                                                                                                       2001US-00993731.
                                                                                                                                                                                                                            2002WO-US035597.
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                                                                                                                                                           ΑT;
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/mod_base= OTHER
/note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                           mod_base= OTHER
                                                      108pp; English
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78.9%;
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Pred. No. 2.1e+02
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tion; inflammation; therapy;
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New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.

heightened

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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappab r; antisense; immune response; infection; tumour; prophylaxis; phosphorothioate; ss.
                                                                                                                                      13-NOV-2001; 2001US-00993731.
                                                                                                                                                            05-NCV-2002; 2002WO-US035597.
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                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                 (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GACCIGCCICTGCAGCIGG
                                                                                         Watt AT;
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/mod_ba
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5-methylcytidines"
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                                                                                                                                                                                                                                 (2'-MOE)
                                                                                                                                                                                                                                                                                                                                     backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              I-kappa-B-related; NFKBIL2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                  nucleotides"
                                                                                                                                                                                                                                                                              nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation; therapy;
                                                                                                                                                                                                                                                                                                                                    All
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                                                                                                                                                                                                                                                                                                                                     cytidine
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Claim 3; Page

108pp;

English

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RESULT 258
AAL61582
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; ikappab r; antisense; immune response; infection; inflammation; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAL61582 standard; DNA; 20
                                                                                                                                                                                                                                       modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
WPI; 2003-468635/44
                           Monia BP,
                                                      (ISIS-) ISIS PHARM INC
                                                                                 13-NOV-2001; 2001US-00993731
                                                                                                           05-NOV-2002; 2002WO-US035597
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Similarity 82.4%;
14; Conservative
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                           Watt AT;
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                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "Phosphorothicate
are 5-methylcytidines"
                                                                                                                                                                                                               /*tag=
/mod_ba
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/mod_base= OTHER
/note= "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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== "2'-methoxyethyl (2'-MOB) nucleotides'
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Pred. No. 2.4e+02;
0; Mismatches 3
                                                                                                                                                                                                                                                        (2'-MOE)
                                                                                                                                                                                                                                                                                                                              backbone; All cytidine residues
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                                                                                                                                                                                                                                                        nucleotides'
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therapy;
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New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened
                                                                                                                                                       immune response or infection.
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Claim 3; Page 75; 108pp; English

The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IXBR, I-kappa-B-related, ikappab r, nuclear factor of kappa BR, polypepides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA

Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Query Match Best Local S Matches 14 ch 0.5%; l Similarity 82.4%; 14; Conservative 0 Score 12.2; DB 1 Pred. No. 2.4e+02 Mismatches DB 1; ω •• Length Indels 0 Gaps 0

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1120

AGGTCCTCCAAGACCTG 1136

밁 AAL61575 standard; DNA; 20 w AGATGCTGCAAGACCTG 19 ВЪ

AAL61575

22-SEP-2003

(first entry)

Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130500.

Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss:

Synthetic. sapiens

modified_base are Location/Qualifiers *tag= base= OTHER

modified_base modified_base 16. .20 /mod_base= OTHER /note= "2'-methoxyethyl *tag= note= "Phosphorothicate backbone; s-methylcytidines" *tag= c base= OTHER (2'-MOE) nucleotides " All cytidine residues

WO2003042360-A2

note= "2'-methoxyethyl

(2'-MOE) nucleotides

22-MAY-2003

05-NOV-2002; 2002WO-US035597.

13-NOV-2001; 2001US-00993731.

Monia BP,

Watt AT;

(ISIS-) ISIS PHARM INC

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RESULT 260
AAL61547
ID AAL615
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Matches 14
                                                                                                                                                                                                                                                                                              Human; inhibitor-kappa B-R; I-kappaBR; ikappab r; antisense; immune response; tumour; prophylaxis; phosphorothicate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened immune response or infection.
                                                                                                                                                                                                                                                                                                                                                         Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130472.
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Synthetic.
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                                                                                                                                               are
                                                                                                                                             /note= "Phosphorothioate backbone;
are 5-methylcytidines"
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                            /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                    /*tag=
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                                                          *tag=
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                                                                                                    mod_base= OTHER
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                                                                                      "2'-methoxyethyl
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Pred. No. 2.4e+02;
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                                                                                                                                                                                                                                                                                                                 IKBR; I-kappa-B-related; NFKBIL2;
infection; inflammation; therapy;
                                                                                                                                                                                                                                                                                                     88
                                                                                         (2'-MOE)
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                                                                                       nucleotides"
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7, 2004, 16:11:53

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GTGTGCGCACAC 20

20

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Gaps

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8
                                                                                                                                                The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I kappaBR, CK INER, I-kappa-B-related, ikappabr, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving crimcreased cytokine expression, or a result of infection (e.g. bacterial, cyiral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide crargetted to human inhibitor-kappa B-R DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heightened
                                                                                                              Sequence
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Conservative (
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                                                             Score 12;
Pred. No.
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                                     DB 1; L.
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US-08-863-639A-71
US-08-863-639A-71
US-08-863-639A-71
US-08-863-639A-71
US-09-863-63-63
US-09-703-63-17
US-09-703-67A-173
US-09-705-267A-173
US-09-705-267A-173
US-09-866-108A-829
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US-09-866-108A-829
US-09-866-108A-8659
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53, Appl
53, Appl
51, Appl
52, Appl
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US-09-866-108A-6391
US-09-866-108A-6658
US-09-866-108A-2615
US-09-866-108A-2616
US-09-866-108A-2616
US-09-866-108A-2616
US-09-866-108A-2618
US-09-866-108A-184
US-08-373-124A-184
US-09-866-108A-2593
US-08-435-628-184
US-09-866-108A-6611
US-09-866-108A-7854
US-09-866-108A-7854
US-09-866-108A-7854
US-09-866-108A-6611
US-
Sequence 668, App Sequence 930, App Sequence 930, App Sequence 2617, App Sequence 6391, App Sequence 6391, App Sequence 6392, App Sequence 6392, App Sequence 184, App Sequence 180, App Sequence 180, App Sequence 184, App Sequence 28, Appl Sequence 28, Appl Sequence 28, Appl Sequence 28, Appl Sequence 187, App Sequence 187, App Sequence 667, App Sequence 107, App Sequence 110, App Sequence 110, App Sequence 110, App Sequence 110, App Sequence 123, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 247, App Sequence 247, App Sequence 247, Appl Sequence 253, Appl Sequence 263, Appl Sequence 263, Appl Sequence 27, Appl Sequence
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                                                                                                                                                                                                                                          TOPOLOGY: linear

MOLECULE TYPE: other nucleic acid

DESCRIPTION: /desc = "Oligonucleotide"
US-08-860-038-18
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Sequence 18, Application US/09580923 Patent No. 6319672 GENERAL INFORMATION:
                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 18: SEQUENCE CHARACTERISTICS: LENGTH: 25 base no:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Re-lease #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/860,038
                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Savitzky Esq., Martin F. REGISTRATION NUMBER: 29,699
REFERENCE/DOCKET NUMBER: ST9408
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION UMBER: FR 94/15162
FILING DATE: 16-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO FR95/01468
APPLICATION NUMBER: WO FR95/01468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, Mailstop 3C43
CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: CROUZET, Joel
APPLICANT: SCHERMAN, Daniel
APPLICANT: WILS, Pierre
TITLE OF INVENTION: PURIFICATION OF A TRIPLE HELIX FORMATION
TITLE OF INVENTION: WITH AN IMMOBILIZED OLIGONUCLEOTIDE
                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 08-NOV-1995
ATTORNEY/AGENT INFORMATION:
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ER: ST94090-US
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US-09-156-979-46
US-09-387-341-107
US-09-866-108A-7854
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Pred. No. 7.
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APPLICANT: Blanche, Francis

APPLICANT: Blanche, Francis

TITLE OF INVENTION: PURIFICATION OF A TRIPLE HELIX FORMATION WITH

TITLE OF INVENTION: IMMOBILIZED OLIGORUCLECTIDE

FILE REFERENCE: 03804.0138-01

CURRENT APPLICATION NUMBER: US/09/580,923

CURRENT FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: 08/860,038

PRIOR APPLICATION NUMBER: 08/860,038

PRIOR APPLICATION NUMBER: PCT/FR95/01468

PRIOR APPLICATION NUMBER: PCT/FR95/01468

PRIOR FILING DATE: 1995-11-08

NUMBER OF SEQ ID NOS: 36

SOPTMARE: Patentin Ver: 2.1

SEQ ID NO 18

LENGTH: 25
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: oligonucleotide
US-09-580-923-18
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Best Local Similarity 95.5%;
Matches 21; Conservative
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GENERAL INFORMATION:
                                                                                                   APPLICATION UNMBER: US/08/863,
APPLICATION MAY 28, 1997
CLASSIFICATION: 435
ATTORNEY, AGENT INFORMATION:
NAME: JOSEPH E. Mueth
REGISTRATION NUMBER: 20,532
REFERENCE, DOCKET NUMBER: 1185
TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 796-4000
                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 in
COMPUTER: IBM compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: COREL Wordberfect 8
CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
                                                                    TELEPHONE: (626) 796-40
TELEFAX: (626) 795-6321
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: OLIGONUCLEOTI
NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheldon & Mak
STREET: 225 South Lake Avenue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Coassin, Peter APPLICANT: Rampal, Jang B. APPLICANT: Caskey, C. T.
                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 91101
                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Pasadena
STATE: CA
STRANDEDNESS:
                               ENGTH:
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            nucleic acid
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Scherman, Daniel
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Coassin, Peter J.
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                                                                                                                                                                                                                                                                                                                                                                 3.50 inch, 1.44 Mb storage
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Pred. No. 7.
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; MOLECULE TYPE:
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                        Query Match
                                                       Sequence 59, Patent No. 59
                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 796-4000
TELEPAX: (626) 795-6321
INFORMATION FOR SEQ ID NO: 53:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Matson
                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                  MOLECULE TYPE: Other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/00
FILING DATE: May 28, 1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IHM COMPUTER: THE CONTROL WINDOWS 95
COTTON DATA: (700 419A)
                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                            STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 20 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                       ENGTH:
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                                                                    Application US/08863639A
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Matson, Robert S. Coassin, Peter J. Rampal, Jang B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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                                                                                                                                                                                              Conservative
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Coassin, Peter J.
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N: 435
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                                                                                                                                                                                                                                                                                         single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jang B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%;
                                                                                                                                                                                                          95.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                              0
                                                                                                                                                                                                          Score 19.4;
Pred. No. 6.
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Pred. No. 6
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                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                      DB 1; Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 64,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
COMPUTER: IBM compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: COTE! WordFerfect 8 version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/863,639A
FILING DATE: May 28, 1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Matson
APPLICANT: Coassi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (626) 795-6321 INFORMATION FOR SEQ ID NO: 59: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH E. Mueth
REGISTRATION UNMER: 20,532
REFERENCE/DOCKET NUMBER: 1185
TELEPHONE: (626) 795-6321
TELEPHAX: (626) 795-6321
                                                                                                                                                                                                                                                                                                                                                                                                                 tent No.
                                                                                                                                   COUNTRY: USA
ZIP: 91101
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheldon &
STREET: 225 South La
                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Caskey, C. T.
TITLE OF INVENTION: OLIGONUCLECTIDE REPEAT ARRAYS
NUMBER OF SEQUENCES: 95
                                                                                                                                                                                                                                                                                           APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
TITLE OF INVENTION: OLIGONUCLECTIDE REPEAT ARRAYS
NUMBER OF SEQUENCES: 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                        STREET: 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
FILING DATE: May 20
CLASSIFICATION: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: Windows 95
SOFTWARE: Corel WordPerfect 8 version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08863639A
                                                                                                                                                                                                                                         225 South Lake Avenue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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25 South Lake Avenue, 9th Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%;
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                                                                                                                                         3.50 inch,
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Pred. No. 6.9;
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                                                                                                                                                                                                                                          9th Floor
                                                                                                                                           1.44 Mb storage
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US-08-863-639A-64

MOLECULE TYPE:

TOPOLOGY: STRANDEDNESS:

linear

single

Other nucleic acid

INFORMATION FOR SEQ ID NO:

(626)

795-6321

64:

ELECOMMUNICATION INFORMATION:

(626)

REFERENCE/DOCKET NUMBER: REGISTRATION NUMBER:

11859-1

ELEPHONE:

SEQUENCE CHARACTERISTICS:

nuc.

21 base pairs ucleic acid

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                                                                                       JS-08-486-421-51
                                           Query Match
Best Local Similarity
                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                         TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO:
                                                                                                    TOPOLOGY: 1
                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/470,911 FILING DATE: 06-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 07-JUN-1
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 30 REFERENCE/DOCKET NUMBER:
                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1770 GAGGAGGAGGAGGAGGAG 1790
   1772 GGAGGAGGAGGCGGAGGAGGCGGC 1795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JCANT: Bergemann, Andrew D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GAGGAGGAGGAGGAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1, Application US/08486421
5672479
                                                                                                                                             I: 24 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                            Coruzzi, Laura A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Johnson, Edward M.
                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pennie & Edmonds
                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-1995
                                                                                                                                                                                                                       869-9741/8864
                                           0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 19.4; DB 1; 95.2%; Pred. No. 6.9; tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                US/08/486,421
                                                                                                                                                                                                                                                                       30,742
                            ; Score 19.2; D; Pred. No. 12; 0; Mismatches
                                                                                                                                                                                                                                                                   6923-053
                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 21;
                                                        Length 24;
                             0
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                          Gaps
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RESULT 9
US-08-486-809-51
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Best Local Similarity 87.5%;
Matches 21; Conservative
                                                                                                                                                                                                                                                                          Sequence 51,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 51,
                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-470-911-51
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                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                APPLICANT: Johnson, Edward M. APPLICANT: Bergemann, Andrew D. TITLE OF INVENTION: CLONING AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAURA A.
REGISTRATION NUMBER: 30,74
REFERENCE/DOCKET NUMBER: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 06-JUN-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10036-2711
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Bergemann, Andrew D.
TITLE OF INVENTION: CLONING AND EXPRESSION OF PUR PROTEIN
                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                           CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
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                                                                                                                             DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                         1772 GGAGGAGGAGGCGGAGGAGGCGGC 1795
                                                                                                                                                                                                                                      INFORMATION:
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                                                                                                                                                                                                                                                                          Application US/08486809
                                                                                                           E: Pennie & Edmonds
1155 Avenue of the Americas
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Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US/08/470,911
                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 19.2;
Pred. No. 12
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                                                                                                                                                                                    EXPRESSION OF PUR PROTEIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 24;
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RESULT 10
US-08-863-639A-70
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Best Local Similarity 87.5%;
Matches 21; Conservative
            COMPUTER: IBM compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Corel WordPerfect 8 version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/863,639A
FILING DATE: May 28, 1997
CLASS.FICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH B. Mueth
REGISTRATION NUMBER: 20,532
REFERENCE/DOCKET NUMBER: 11859-1
TELECOMMUNICATION: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Matson, Robert S.
APPLICANT: Coassin, Peter J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                    ZIP: 91101
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA
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FILING DATE: 07-UTN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/470,911
PRIOR APPLICATION NUMBER: US 08/470,911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: COTUZZI, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 692:
RELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 06-JUN-1995
TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                  COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Sheldon & ran
STREET: 225 South Lake Avenue, 9th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                   RRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
CELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               E OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1772 GGAGGAGGAGGCGGAGGAGGCGGC 1795
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                                                                                                                                                                                                                                                                                                                        USA
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VENTION: OLIGONUCLEOTIDE REPEAT ARRAYS
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  (626)
                                                                                                                                                                                                                                                          Diskette, 3.50 inch, 1.44 Mb storage
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796-4000
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Pred. No. 12;
O; Mismatches
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US-08-863-639A-83/c
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                                                                                                                                                                                                                                                                                   DIBKETTE, 3.50 inch, 1.44

OPERATING SYSTEM: Windows 95

SOFTWARE: Corel Wordberfect 8 version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/863,639A

FILING DATE: May 28, 1997
CLASSIFICATION: 435
ATTORNEY/AGERM
ATTORNEY/AGERM
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Best Local Simi
Matches 19;
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                                                 Query Match
Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                 TELEFAX: (626) 795-6321
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                        TELEPHONE: (626) 796-4000
                                                                                                                   MOLECULE TYPE: Other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Matson, Robert S. APPLICANT: Coassin, Peter J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Coassin, Peter APPLICANT: Rampal, Jang B.
                                                                                                                                                                                                                                                                 NAME: JOSEPH E. Mueth
REGISTRATION NUMBER:
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                                                                                                                                 STRANDEDNESS:
TOPOLOGY: li
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                                                                                                                                                                       LENGTH:
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                         1772 GGAGGAGGAGGCGGAGGAGG 1791
21
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                                                   0.7%; Score 18.4; ilarity 95.0%; Pred. No. 1 Conservative 0; Mismatch
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JENTION: OLIGONUCLECTIDE REPEAT ARRAYS
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                                                                                                                                         single
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Pred. No. 1:
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RESULT 12 US-08-486-343A-5/c

Sequence 5, Application US/08486343A Patent No. 6071695

APPLICANT:

OZKAYNAK, ENGIN

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RESULT 13
PCT-US95-07349-5/c
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                                                                                                                                                                                                                                                                                          Sequence 5, Application PC/TUS9507349 GENERAL INFORMATION:
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Best Local Similarity
Matches 19; Conserv
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: floppy disk
COMPUTER: IEM PC COMPATIBLE
COMPUTER: IEM PC COMPATIBLE
COMPUTER: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/486,343A
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                        TITLE OF INVENTION: METITLE OF INVENTION: MCNUMBER OF SEQUENCES: 7
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REFERENCE/DOCKET NUMBER: CR.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)-248-7000
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LOCATION: 1..21
OTHER INFORMATION:
OTHER INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
RRENT APPLICATION DATA:
                                                                                               COUNTRY: U
                                                                                                                                           CITY: HOPKINTON
                                                                                                                                                            STREET:
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                             METHODS AND COMPOSITIONS FOR MODULATING MORPHOGEN EXPRESSION
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Pred. No. 11
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                   Version #1.25
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US-09-433-699-43/c
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US-09-490-692-153/c
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LOCATION: 1..21
OTHER INFORMATION: /note= "WT1 HUMAN TCC BINDING SITE"
PCT-US95-07349-5
Sequence 153, Application US/09490692
Patent NO. 6180353
GENERAL INFORMATION:
APPLICANT: Nicholas M. Dean
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF DAXX EXPRESSION
FILE REFERENCE: RTS-0120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 43
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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GENERAL INFORMATION:
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Best Local Similarity 94.7
Matches 18; Conservative
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Best Local Similarity
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APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTIGENSE MODULATION OF NUCLEOLIN EXPRESSION
FILE REFERENCE: RTS-0109
FOURRENT APPLICATION NUMBER: US/09/433,699B
CURRENT FILING DATE: 1999-11-03
RUMBER OF SEQ ID NOS: 89
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LENGTH: 21 base pairs
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TORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ECOMMUNICATION INFORMATION:
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Pred. No. 11
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Pred. No. 1
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-444-053-54
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Patent No. 6165728
GENERAL INFORMATION:
APPLICANT: Donna T. Ward
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF NCK-2 EXPRESSION
FILE REFERENCE: RTS-01.22
CURRENT APPLICATION NUMBER: US/09/444,053A
CURRENT FILING DATE: 199-11-19
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 54
LENGTH: 20
TYPE-. NO.
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Best Local Similarity
Matches 18; Conserv
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Sequence 12, App-
No. 615971
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CURRENT FILING DATE: 2000-01-24
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EQ ID NO 153
LENGTH: 20
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                      ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Cntr., P.O. Box 457
CITY: Spring House
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                        COUNTRY:
                                                                                                                                                                                                                                 LICANT: Krummenacher, Claude F.

LE OF INVENTION: Method and Compositions for Stabilizing

LE OF INVENTION: Unstable Gene Transcripts
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                                                                                                                                                                                                                                                                                                                                             INFORMATION:
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Pred. No. 14;
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Pred. No. 18
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                                                                                                                                      US-09-733-444-22
                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 22
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
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Matches
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                                                                    Matches
                                                                                                   Query Match
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                        APPLICANT: University of Nebraska Board of Regents TITLE OF INVENTION: Specific Mucin Expression as a NTITLE OF INVENTION: for Pancreatic Cancer FILE REFERENCE: UNMC 63155 CURRENT APPLICATION NUMBER: US/09/733,444 CURRENT APPLICATION NUMBER: US/09/733,444 CURRENT FILING DATE: 2000-12-08
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Batra, Surinder K.
APPLICANT: Brandt, Randall E.
APPLICANT: Ringel, J"erg
                                                                                                                                                                         FEATURE:
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DESCRIPTION: /desc = "probe/primer Exon
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ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER: WST78APCT
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                                  894 CTGCAGCAGACAGCCCTG 911
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5, 6576423
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18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Faulmann, Grit
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                                                                     Conservative
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90.0%; Pred. No. 21;
tive 0; Mismatches
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                                                                                  Score 16.4;
Pred. No. 1
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                                                                                                   Length: 18;
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US-09-780-173A-93; Sequence 93, Application US/09780173A; Patent No. 6455307

RESULT 19

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RESULT 20
US-09-490-692-155/c
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                                                                                                                                                                                                                                                                        US-09-705-267A-173
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Best Local Similarity
Matches 17; Conserv
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TYPB: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
3-09-780-173A-93
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SEQ ID NO 93
                                             NUMBER OF SEQ ID NOS:
SEQ ID NO 173
LENGTH: 20
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GENERAL INFORMATION:
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Patent No. 6551826
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APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION; ANTISENSE MODULATION OF CASEIN KINASE 2-ALPHA PRIME EXPRESSION
FILE REFERENCE: RTS-0165
CURRENT APPLICATION NUMBER: US/09/780,173A
CURRENT FILING DATE: 2001-02-08
                                                                                                                                                                                                                     GENERAL INFORMATION:
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CURRENT FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 176
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APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF DAXX EXPRESSION
FILE REFERENCE: RTS-0120
                                                                                              APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF RAIDD EXPRESSION
FILE REFERENCE: RTS-0211
CURRENT APPLICATION NUMBER: US/09/705,267A
CURRENT FILING DATE: 2000-11-01
                                                                                                                                                                                                   APPLICANT: Hong Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Pred. No. 21;
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US-09-705-267A-174
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Best Local S
Matches 16
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Patent No. 5561041
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.
CURRENT APPLICATION NUMBER: US/08/152,313
APPLICATION NUMBER: US/08/152,313
FILING DATE: 12-NOV-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, Jr., Ph.D., John R.,
REGISTRATION NUMBER: PD-2912
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF RAIDD EXPRESSION
FILE REFERENCE: RTS-0211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/705,267A
CURRENT FILING DATE: 2000-11-01
NUMBER OF SEQ ID NOS: 177
                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 20
TYPE: DNA
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                                                                                                                                                                                                                                                             ADDRESSEE: Spensley Horn Jubas & Lubitz
STREET: 1880 Century Park East, Suite 500
CITY: Los Angeles
STATE: California
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les 16; Conservative 0;
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5. 6551826
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ANALYSIS OF SPUTUM
128
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.00.0%; Pred. No.
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Pred. No.
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US-08-152-313-113
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Best Local Similarity 94. Matches 16; Conservative
                                  Query Match
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INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                         TELEFAX: (619) 455-5110
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                  FILING DATE: 12-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, Jr., Ph.D., John
REGISTRATION NUMBER: 31,678
REFERENCE/DOCKET NUMBER: PD-2912
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/579,223
FILING DATE: 28-DEC-1995
                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 455-5100
                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/152,313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sidransky, David
TITLE OF INVENTION: NUCLEIC ACID MUTATION DETECTION BY
TITLE OF INVENTION: ANALYSIS OF SPUTUM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic)
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LOCATION: 1...
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TOPOLOGY: linear
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5. 5726019
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1880 Century Park East, Suite 500
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0.6%; Score 15.4; I
94.1%; Pred. No. 20;
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94.1%;
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Pred. No. 20
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Gaps
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SOFTWARE: Aeomica Sequence
Patent No. 6686188
SEO ID NO 929
LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
US-09-866-108A-929
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: ABOMICA-7
FULE REFERENCE: ABOMICA-7
CURRENT APPLICATION UNMBER: US/09/866,108A
CURRENT FILLING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
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US-09-866-108A-8659
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US-09-866-108A-929
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GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 15755
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PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
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                                                                                                                                                                 INFORMATION:
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o. 6686188
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RANK, David R.
CHEN, Wensheng
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Pred. No. 2
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        HEART AND MUSCLE
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RESULT 27
PCT-US94-12947A-113/c
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; ORGANISM: Homo sapiens 
US-09-866-108A-8659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Aeomica Sequence Listing Engine Patent No. 6686188
SEQ ID NO 8659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2001-01-30
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
ATTORNEY/AGENT INFORMATION:
NAME: Haile, Ph.D., Lisa A.
REGISTRATION NUMBER: P-38,347
REFERENCE/DOCKET NUMBER: FD-2:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 455-5100
                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                       STREET: Los Angeles
CITY: Los Angeles
STATE: California
                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: The Johns Hopkins University School of Medicine TITLE OF INVENTION: NUCLEIC ACID MUTATION DETECTION BY TITLE OF INVENTION: ANALYSIS OF SPUTUM
                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                           APPLICATION NUMBER: FILING DATE: 10-NOV CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                     COUNTRY: U
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US01/00663
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FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                 3: Spensley Horn Jubas & Lubitz
1880 Century Park East, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application PC/TUS9412947A
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                                                                                                                                               10-NOV-1994
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                                                                                                                                                                 PCT/US94/12947A
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                                          FD-2912
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Pred. No. 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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US-09-387-341-107/c
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PCT-US94-12947A-113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-156-979-46/c
                                                                                                                                                                     APPLICANT: Roberts, M. Luisa
APPLICANT: Cowsert, Lex M.
TITLE OF INVENTION: Antisense Modulation of
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0404
CURRENT APPLICATION NUMBER: US/09/387,341
CURRENT FILING DATE: 1999-08-31
CURRENT APPLICATION NUMBER: 09/156,424
FERRITER APPLICATION NUMBER: 09/156,424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 46
LENGTH: 18
SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Cowsert, Lex M.
TITLE OF INVENTION: ANTISENSE MODULATION OF Rhob EXPRESSION
                                                                                                                                                                                                                                                                                                                                                   Sequence 107, Application US/09387341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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Best Local Similarity
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Patent No. 596267
                                                                                                                                                                                                                                                                                                                                  Patent No. 64103
                                             EARLIER APPLICATION NUMBER: 09/156,979
EARLIER FILING DATE: 1998-09-18
EARLIER APPLICATION NUMBER: 09/156,807
EARLIER FILING DATE: 1998-09-18
EARLIER APPLICATION NUMBER: 09/161,015
EARLIER FILING DATE: 1998-09-25
NUMBER OF TEXTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: RTS-0013
CURRENT APPLICATION NUMBER: US/09/156,979
CURRENT FILING DATE: 1998-09-18
NUMBER OF SEQ ID NOS: 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                                    INFORMATION:
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                                                                                                                                                          FILING DATE: 1998-09-18
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                PatentIn Ver.
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 Mismatches

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Pred. No. 30;
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Pred. No. 2
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; OTHER INFORMATION: Description of Artificial Sequence:Synthetic US-09-387-341-107

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<u>,,</u>

Length 18;

Query Match

FEATURE

TYPE: DNA
ORGANISM: Artificial Sequence

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                                                                                                                         JS-08-584-040-5442
                                                                                                                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                  Query Match 0.6%;
Best Local Similarity 62.5%;
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/584,0
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCCODET 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: ESCOBEDO, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHBLIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
REFERENCE/CATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
ADDRESSEE: Lyon & Ly
                                                                                                                                    STRANDEDNESS: 811
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A. ZIP: 90071-2066
1301 CATGGTCATCTGTGAG 1316
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o. 6346398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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tinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pamela
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                                                  Score 14.4; DI
Pred. No. 30;
5; Mismatches
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Pred. No. 3
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                                                                                      <u>,</u>
                                                     Indels
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                                                     Gaps
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US-09-920-663-4/c
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                                                                                                                       TYPE: RNA
ORGANISM: Homo sapiens
US-09-474-432B-669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/09920663 Patent No. 6426221 GENERAL INFORMATION:
                                                                                                                                                                                    SEQ ID NO 669
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 49
SEQ ID NO 4
                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                              APPLICANT: Sweedler, David
APPLICANT: Sweedler, David
APPLICANT: Zinnen, Shawn
ITTLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonucleoti
ETLE REFERENCE: MBHB00-831-B (247/276)
ETLE REFERENCE: MBHB00-831-B (247/276)
ETLE REFERENCE: MBHB00-831-B (247/276)
CURRENT FILING DATE: 1999-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR PILING DATE: 1998-04-29
PRIOR PILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/301,511
PRIOR APPLICATION NUMBER: US 09/301,511
PRIOR FILING DATE: 1999-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -09-474-432B-669/c
                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 1526
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/920,663
CURRENT FILING DATE: 2001-08-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Donna T. Ward
APPLICANT: Lex M. COWSETT
TITLE OF INVENTION: ANTISENSE MODULATION OF RIP2 EXPRESSION
FILE REFERENCE: RTS-0233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Beigelman, Leo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 17
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                               Local
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    1266 CTGGAAGAGGCTGAGG 1281
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15; Conservative
                                         l Similarity
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Adamic, Jasenka
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                                           Conservative
                                                            93.88;
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93.8%; Pred. No. 3
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                                                               Score 14.4;
Pred. No. 3
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                                                                                DB 1;
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US-09-476-387-668/c
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PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/301,511
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR PPLICATION NUMBER: 60/083,727
PRIOR PILING DATE: 1998-14-29
PRIOR APPLICATION NUMBER: 60/064,866
PRIOR FILING DATE: 1997-11-05
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Query Match
Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleot FILE REFERENCE: MBHB00-831-C (249/073)
CURRENT APPLICATION NUMBER: US/09/476,387
CURRENT FILING DATE: 2001-04-04
PRIOR ADDITATION TOTAL CONTROL TO THE TOTAL CONTROL TOTAL 
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
APPLICANTON: Levels of Vascular Endothelial Growth Factor Receptor
FILE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT FILING DATE: 1996-01-08
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
NUMBER: OF SEO ID NOS: 14225
COMMUNER OF SEO ID NOS: 14225
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APPLICANT: Beigelman, Leo
APPLICANT: Beaudry, Amber
APPLICANT: Karpeisky, Alex
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                                                                                                                                                                                                                                                                                                                                    TYPE: RNA
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0.6%;
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              Score 14.4;
Pred. No. 30;
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Pred. No. 3
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; TYPE: DNA
; ORGANISM: Homo
US-09-866-108A-928
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US-09-866-108A-930
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PRIOR FILING DATE: 2000-10-04
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
                                                                                                                                                                    Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                         Patent No. 66
SEQ ID NO 928
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
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APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
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                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Aeomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                                                                                                                                      Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
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                                                                                                                                                                                                                                                                                                                    LENGTH: 17
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APPLICATION NUMBER: PCT/US01/00663
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                                                                                                                            1264 AGCTGGAAGAGGCTGA 1279
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5. 6686188
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HANZEL, David K.
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Sequence 930, Application US/09866108A Patent No. 6686188

INFORMATION:

APPLICANT: GU, Yizhong APPLICANT: JI, Yonggang APPLICANT: PENN, Sharron

HANZEL, David K RANK, David R. CHEN, Wensheng

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Remaining Prior Application d.
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Li:
Patent No. 6686188
SEQ ID NO 930
LENGTH: 17
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US-09-866-108A-930
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Best Local
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
CURRENT APPLICATION NUMBER: US 60/207,456
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PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: US 60/236,359
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
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                                                          OR FILING DATE: 2000-05-26
OR APPLICATION NUMBER: GB 24263.6
OR FILING DATE: 2000-10-04
OR APPLICATION NUMBER: US 60/236,359
OR FILING DATE: 2000-09-27
OR APPLICATION NUMBER: PCT/US01/00666
OR APPLICATION NUMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
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PLICATION NUMBER: PCT/US01/00669
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CATION NUMBER: PCT/US01/00663
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RANK, David R.
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; ORGANISM: Homo sapiens US-09-866-108A-2618
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SEQ ID NO 2617
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
                                                                                                              NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Aeomica Sequence Listing
                                                                                                                                                                       PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR FILING DATE: 2000-10-04
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
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NUMBER OF SEQ ID NOS: 15755
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PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
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SEQ ID NO 6391
                                                                                                Sequence 6392, Application US/09866108A Patent No. 6686188
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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TYPE: DNA
ORGANISM: Homo sapiens
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                                                   APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
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                                                                                INFORMATION:
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JI, Yonggang
PENN, Sharron G
HANZEL, David K
RANK, David R.
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PENN, Sharron G.
HANZEL, David K.
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FILE REFERENCE: ABOMICA-7;
FILE REFERENCE: ABOMICA-7;
CURRENT PILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR APPLICATION NUMBER: DCT/US01/00666
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SEQ ID NO 6392
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PRIOR PILLING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILLING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILLING DATE: 2000-09-27
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APPLICANT: JI, Yongga
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TYPE: DNA
ORGANISM: Homo sapiens
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RPPLICANT: SHANNON, Mark
RITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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TLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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FILING DATE: 2001-01-30
APPLICATION NUMBER: ECT/US01/00665
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
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HANZEL, David
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667

2001-01-30

LING DATE:

: 2001-01-30 NUMBER: PCT/US01/00664 : 2001-01-30

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; ORGANISM: Homo sapiens
US-09-866-108A-8658
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Best Local Similarity 93.8%;
Matches 15; Conservative
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SEQ ID NO 8658
LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
-09-866-108A-8660
                                                                                       SEQ ID NO 8660
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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                                                                                                                               SOFTWARE: Aeomica
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
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OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/USO1/00667
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/USO1/00664
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/USO1/00669
OR APPLICATION NUMBER: PCT/USO1/00669
OR APPLICATION NUMBER: PCT/USO1/00665
OR APPLICATION NUMBER: PCT/USO1/00665
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APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
                                                                                                                                                                                                     FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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F: SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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ING DATE: 2001-01-30
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HANZEL, David K.
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                                                                                                                                                                              Application data removed -
                                                                                                                                   Sequence Listing Engine
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Pred. No. 3
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                                                                                                                                                                                See File Wrapper or PALM
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APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert

TITLE OF INVENTION: ANTISENSE MODULATION OF SMAD2 EXPRESSION
FILE REFERENCE: RTS-0044
CURRENT APPLICATION NUMBER: US/09/255,912
CURRENT FILING DATE: 1999-02-23
NUMBER OF SEQ ID NOS: 47
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/09255912 Patent No. 6037142
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Best Local
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                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/866,108A CURRENT FILING DATE: 2001-05-25 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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TLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                                                                                              APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
                                                                                                                                                                  FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/236,359
                                                             APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
                                                                                                  FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                          APPLICATION NUMBER: GB 24263.6
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                                                APPLICATION NUMBER: PCT/US01/00665
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5. 6686188
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Similarity 93.8%;
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             G DATE: 2001-01-30
CATION NUMBER: PCT/US01/00668
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93.8%; Pred. No. 3
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Pred. No. 3
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Best Local Similarity
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Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
                                                                                                                                                                                                                           SOFTWARE: Aeomica Sequence Listing Engine
                                                                                                                                                                                                                                        Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: GB 24263.6
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                               1840 TCTCAGAGAGCGAG 1853
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3. 6686188
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                                                                              Similarity
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CATION NUMBER: PCT/US01/00669
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ION NUMBER: US 60/207,456
ATE: 2000-05-26
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N, Sharron G.
NZEL, David K.
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100.0%; Pred. No. 35;
tive 0; Mismatches
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RESULT 46
US-08-373-124A-178/c
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                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 17 barr
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                                                   Matches
                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: January PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                    TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                   FILING DATE: August 26, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                    TELEPHONE: (213)
                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.O
                                                              Local
                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                    FILING DATE: Februa
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
                                                                                                                                                                                                                                                            AME: Warburg, Richa
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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                        1774 AGGAGGAGGCGGAGGAG 1790
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  17
                                                 l Similarity 88.2
15; Conservative
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California
  AGGAGAAGGAGGAGGAG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      633 West Fifth Street
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                                                                                                                s: single
linear
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                                                                                                                                                                                                                                                                                                                                                                                                                   January 13, 1995
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                                                               88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                         08/245,466
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                                                                                                                                                                                                                                                                                                                                                     1994
                                                 0
                                                                                                                                                                                                                                                 209/035
                                                                            Score 13.8;
                                                   Pred. No. 38;
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOS 5.0
                                                                            DB 1;
                                                    2
                                                                          Length 17;
                                                    Indels
                                                    0
                                                    Gaps
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US-08-373-124A-180/c ; Sequence 180, Application US/08373124A

0

Patent No.

SENERAL INFORMATION:

APPLICANT:

Stinchcomb, Dan T. Draper, Kenneth

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RESULT 48
US-08-373-124A-182/c
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TELEPAX: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 180:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 Dase pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/373,124A
FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/986,422
PILING DATE: AUGUST 26,1992
APPLICATION NUMBER: 07/936,422
                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                             tent No.
                                                                           APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Warburg, Richard REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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STATE: California
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o. 5646042
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LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 184, Application US/08373124A Patent No. 5646042
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FILING DATE: January 13, 1995
PRIOR APPLICATION DATA: 08/245,466
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Stinch
APPLICANT: Draper
                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                           COUNTRY: U.S.A.
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Jarvis, Thale
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บร-08-261-822A-30/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: May 18, 194
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,32
FILING DATE: February 7, 1992
APPLICATION NUMBER: 07/986,422
APPLICATION NUMBER: 07/936,422
FILING DATE: August 26, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 299/035
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                atenc No. The Applicant:

GENERAL INFORMATION:

APPLICANT: Ecker, Joseph R. et al.

APPLICANT: THE APPLICANT SERVICE PLANT Genes for Sensitivity to Ethylene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/261,822A
FILLING DATE: 17-UN-1994
CLASSIFICATION: 536
CUARSIFICATION: 536
        ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: January 13, 1995,
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (2-1)
TELEFAX: 67-3510
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nes 15; Conserv
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                                                                                                                                                                                                                                                                                                        COUNTRY: U
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USA
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we Liberty Place, 46th floor
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568-3439
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88.2%; Pred. No. 38;
ative 0; Mismatches
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              TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                               APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
APTLING DATE: AUGUST 26, 1992
APTORNEY/AGENT INFORMATION:
                                                                                              NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 88.2%;
                                                                               TELEPHONE: (213) 489-161
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Los Angeles
: California
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5817796
nucleic acid
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633 West Fifth Street
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TV 13, 1995
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                                                                                                                                      209/035
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Pred. No. 3
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US-08-435-628-180
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                                                                                                                    NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 180:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
Query Match 0.5%; Score 13.8; I Best Local Similarity 88.2%; Pred. No. 38; Matches 15; Conservative 0; Mismatches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILLING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILLING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILLING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILLING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILLING DATE: August 26, 1992
APPLICATION NUMBER: 07/986,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM COMPATIBLE
COMPUTER: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND TITLE OF INVENTION: CANCER USING RIBOZYMES OF SEQUENCES: 2627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 05-MAY CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Suite 4700
CITY: Los Angeles
STATE: California
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5817796
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633 West Fifth Street
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McSwiggen, James
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05-MAY-1995
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Pred. No. 3
                                             DB 1; Length 17;
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                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                      Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/887,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/966,422
FILING DATE: December 26, 1992
APPLICATION NUMBER: 07/966,422
FILING DATE: August 26, 1992
ATTORNEY/AGRET INFORMATION:
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FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 209
                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM COMPONERATING SYSTEM:
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ZIP: 9007
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CITY: Los Angeles
STATE: California
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                                                                                                                                                          TOPOLOGY:
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17 GAGGAGGAGGAGGA 1
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                                                                                       Similarity
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633 West Fifth Street
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                                                                        Conservative
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                                                                                                                                                            linear
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                                                                                                                                                                          single
                                                                    Score 13.8; D
Pred. No. 38;
0; Mismatches
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                                                                                                         DB 1;
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                                                                                                       Length 17;
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RESULT 54

US-08-435-628-184/c

184, Application US/08435628 5. 5817796

INFORMATION:

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                                                                                                            RESULT 55
                                                                                        US-08-613-417A-28
                                                                                                                                                                                                                                                                                                                   US-08-435-628-184
                                     Sequence 28, Application US/08613417A Patent.No. 5874553
GENERAL INFORMATION:
                                                                                                                                                                                                                                        Query Match 0.5%;
Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: August 26, 1992
FILING DATE: August 26, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Stingnow,
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHONS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 67-3510 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS LENGTH: 17 base pairs Type: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C.
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Warburg, Richard REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 20: TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICANT:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: si:
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
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STREET: 633 West Fifth Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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: California
'RY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER: 08/192,943
NUMBER: 7, 1994
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   Phosphonomonoester nucleic acids,
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                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                          Score 13.8;
Pred. No. 3
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FILING DATE:
CLASSIFICATION: 514
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 15; Conserv
                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 195 02 912.7
PRIOR DATE: 31-0AN-1995
ATTORNEY,AGENT INFORMATION:
NAME: SANDERCOCK, Colin G.
REGISTRATION NUMBER: 31,298
REFERENCE/DOCKET NUMBER: 18748/264/HOCE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202672-5300
TELEX: 904136
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tent No.
                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/594,452
FILLING DATE: 31-JAN-1996
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                       ZIP: 20007-5109

COMPUTER READABLE FORM:
MEDIUM TYPE; Floppy disk
COMPUTER: ISM PC compatible
COMPUTER: ISM PC COMPATIBLE FOLIOS/NS-DOS
CORRESTING SYSTEM: PC-DOS/NS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: PEYMAN, Anu APPLICANT: UHLMANN, EU APPLICANT: UHLMANN, EU APPLICANT: OF TNVENTION: G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IMP C Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0. Version #1.25 (EPO)
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: Duc.
STRANDEDNESS: SI._
TOPOLOGY: linear
TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 3000 K SI
CITY: Washington
                                   ELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
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                                      (202) 672-5399
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G CAP-STABILIZED OLIGONUCLEOTIDES
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Pred. No. 3
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QUENCE CHARACTERISTICS: LENGTH: 17 base pairs

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RESULT 58
US-09-196-132-28
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US-09-258-408-28
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; TOPOLOGY: lir
US-08-594-452-28
                                                                                                                                                                                                                                                                                          JS-09-258-408-28
                              Sequence 28, Application US/09196132 Patent No. 6127346 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 28, Application US/09258408 Patent No. 6121434
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Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MBDIUM TYPE: Floppy disk
COMPUTER: IBM FOC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 31,298
REFERENCE/DOCKET NUMBER: 18748/264/HOCE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: PEYMAN, AUGUSTA PELICANT: UHLMANN, EUGEN
TITLE OF INVENTION: G CAP-STABILIZED OLIGONUCLEOTIDES
TOTAL TOTAL PROPERCIES: 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTORNEY/AGENT INFORMATION: NAME: SANDERCOCK, Colin
ITLE OF INVENTION:
                                                                                                                                                                                                              0.5%;
Local Similarity 88.2%;
es 15; Conservarion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/258,408 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (202) 672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
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Phosphonomonoester nucleic acids,
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Pred. No. 3
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Pred. No. 3
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                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-196-132-28
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                                                               CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                            CURRENT APPLICATION DATA APPLICATION NUMBER: US FILING DATE: January:
                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0. Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/196,132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY:
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MOLECULE TYPE:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 8
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APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
TTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 88.2
les 15; Conservative
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5. 6346398
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REMCSWIGGEN, James

Stinchcomb, Dan T.

T: Stinchcomb, Jaime
T: Escobedo, Jaime
TI: Escobedo
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633 West Fifth Street
Suite 4700
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                                                                                                                                                         UMBER: US/08/584,040
January 11, 1996
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OF VASCULAR ENDOTHELIAL
GROWTH FACTOR
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Pred. No. 38;
0; Mismatches
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Warburg,

Richard J

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TELEX: (214)
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
SEQUENCE: 17 base pairs
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                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 8torage
COMPUTER: IBM Compatable
COMPUTER: IBM Compatable
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
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LENGTH: 17 base pairs
TYPE: nucleic acid
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REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                          NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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STREET: Suite 4700
CITY: Los 777
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TOPOLOGY: lir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                          TELEPHONE: (213) 489-16
TELEFAX: (213) 955-0440
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: Los Angeles
S: California
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Stinchcomb, Dan T.
Stinchcomb, Jaine
Escobedo, Jaine
Evention: METHOD AND REAGENT FOR THE
TVENTION: TREATMENT OF DISEASES OR
TVENTION: CONDITIONS RELATED TO LEVELS
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                                                                                                                                                 218/064
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Pred. No. 3
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US-08-584-040-5441
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Robozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaine
TITLE OF INVENTION: Levels of Vascular Endot
FILE REFERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION UNMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION UNMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
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TITLE OF INVENTION: Nucleotide triphosphate a
FILE REFERENCE: MBHBOO-831-B (247/276)
CURRENT APPLICATION NUMBER: US(09/474,432B
CURRENT FILING DATE: 199-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR APPLICATION NUMBER: US 60/064,727
PRIOR FILING DATE: 1997-11-05
PRIOR FILING DATE: 1998-04-29
PRIOR PPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/301,511
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Matches 10; Conserv
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 668
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                                                                                                                                                                                                                                                                                                                           Sequence 1607, Application US/09371772B Patent No. 6566127
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APPLICANT:
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Adamic, Jasenka
Sweedler, David
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Pred. No. 38;
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247/276)
                                                                                                                                                Endothelial Growth Factor Receptor
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PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/301,511
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1998-04-29
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Best Local
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Best Local Similarity
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-09-476-387-667
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CURRENT APPLICATION NUMBER: US/09/476,387
CURRENT FILING DATE: 2001-04-04
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SOFTWARE: PatentIn version 3.0
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TLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleot
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ENGTH: 17
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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Beaudry, Amber
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ilarity 88.2%;
Conservative
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                                                               Yonggang
NN, Sharron G.
NZEL, David K.
              Wensheng
                                                  David R.
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Pred. No. 38;
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Pred. No. 3
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                                                                 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILLING DATE: 2000-10-04
PRIOR PEPLICATION NUMBER: US 60/236,359
PRIOR PEPLICATION NUMBER: US 60/236,359
PRIOR PEPLICATION NUMBER: PCT/USO1/00666
PRIOR PEPLICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR PRICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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Best Local Similarity
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                                CATION NUMBER: PCT/US01/00669
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PCT/US01/00665
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Pred. No. 38;
O; Mismatches
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 15755
PRIOR FILING DATE: 15755
SOFTWARE: Aeomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 2593
LENGTH: 17
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Query Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                 Patent No. 6686188
SEQ ID NO 6611
LENGTH: 17
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 2000-09-27
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
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                                                                                                                                                                                             SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                                                                                             Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
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                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
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TILE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
TILE REFERENCE: ABOMICA-7
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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PENN, Sharron G.
HANZEL, David K.
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                      88.2%;
      0
                      Score 13.8;
Pred. No. 38;
        Mismatches
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RESULT 67
US-09-866-108A-6612/c
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US-09-866-108A-6612
RESULT 68
US-09-866-108A-7854/c
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 6612
LENGTH: 17
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                                                                                                                                                                        Query Match
Best Local Similarity
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: ADMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
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PRIOR FILING DATE: 2000-10-04
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00663
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SL, David K.
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Pred. No. 3
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GENERAL INFORMATION:
APPLICANT: GU, Yizhong

JI, Yonggang PENN, Sharron

David

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SHANNON, Mark RANK, David R. CHEN, Wensheng HANZEL, Sequence 7854, Application US/09866108A Patent No. 6686188

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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 686188
SEQ ID NO 7854
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Best Local &
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TITLE OF II
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PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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                                                        OR FILING DATE: 2000-10-04
OR APPLICATION NUMBER: US 60/236,359
OR FILING DATE: 2000-09-27
OR APPLICATION NUMBER: PCT/US01/00666
OR FILING DATE: 2001-01-30
OR APPLICATION UMBER: PCT/US01/00667
OR APPLICATION UMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
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nes 15; Conserv
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APPLICATION NUMBER: US 60/207,456
FILING DATE: 2000-05-26
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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                                APPLICATION NUMBER: PCT/US01/00664
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LING DATE: 2001-01-30
PLICATION NUMBER: PCT/US01/00664
LING DATE: 2001-01-30
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
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ON NUMBER: PCT/US01/00669
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; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108A-8082
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108A-7855
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                                                                                                                             SOFTWARE: Aeomica Sequence Listing Engine Patent No. 6686188
SEQ ID NO 8082
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Patent No. 6686188
SEQ ID NO 7855
LENGTH: 17
Query Match
Best Local Similarity
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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CURRENT FILING DATE: 2001-05-25
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION UNMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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al Similarity 88.2%;
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INVENTION: MYOSIN-LIKE GENE EXPRESSED
ERENCE: AEOMICA-7
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CATION NUMBER: PCT/US01/00663
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FNN, Sharron G.
Tr., David K.
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Pred. No. 3
  Score 13.8;
Pred. No. 3
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                  Length 17;
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Best Local :
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SEQ ID NO 8648
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
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APPLICATION NUMBER: PCT/US01/00663
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                                                                                                                                                                                                                                1254 CTGCAGCAACAGCTGGA 1270
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T: CHEN, Wensheng
T: SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED
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                                                                                                                                                                                                   CTGCAGCTGCAGCTGGA 17
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G DATE: 2000-09-27
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                               HANZEL,
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ilarity 88.2%;
Conservative
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                                              Yonggang
N, Sharron
David K.

David R.

Wensh
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Pred. No. 38;
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6886188
SEQ ID NO 10738
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 10740, Application US/09866108A Patent No. 6686188
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Best Local Similarity
Matches 15; Conserv
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
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PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
                                                                                                                                                                                                                                                                                                                                                                                                        ENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga:
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                                                                                   PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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TLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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APPLICATION NUMBER: PCT/US01/00667
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00663
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                                                                     ICATION NUMBER: PCT/US01/00667
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HANZEL, David
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                                   NUMBER: PCT/US01/00664
NUMBER: PCT/US01/00669
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Pred. No. 38
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US-09-866-108A-10740
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; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR PRIOR DATE: 2001-01-30
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; Remaining Prior Application data removed - Se NUMBER OF SEQ ID NOS: 15755
                                                                             ; MOLECULE TYPE: (
; HYPOTHETICAL: NO
; ANTI-SENSE: YES
PCT-US95-07744A-30
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SEQ ID NO 10740
LENGTH: 17
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                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/261,8:
FILING DATE: June 17, 1994
ATTORNEY AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: DE PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Relaction DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Trustees of The University of Pennsylvania
TITLE OF INVENTION: Plant Genes for Sensitivity to Ethylene
TITLE OF INVENTION: and Pathogens
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                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                         TOPOLOGY:
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ch 0.5%; Score 13.8; I Similarity 88.2%; Pred. No. 38; 15; Conservative 0; Mismatches
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                                                                                                                                                                                                               17 base pairs
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One Liberty Place, 46th floor
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                                                                                                                                                                                                                                                                            (215) 568-3439
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88.2%; Pred. No. 38;
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                                         DB 1; Length 17;
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TELEFAX: 212-664-0525
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genom
US-08-242-664-19
                                                                RESULT 76
US-08-484-138-19
; Sequence 19, Ap;
; Patent No. 5652
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Matches
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 GENERAL INFORMATION:
APPLICANT: Watanal
APPLICANT: Ren, W
APPLICANT: Weil,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION: APPLICANT: Watana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/242,66
FILING DATE: May 12, 1994
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CITY: New York
STATE: New York
COUNTRY: U.S.A.
                                                                                                                                                                                                                     y match 0.5%; Score 13.4; Local Similarity 93.3%; Pred. No. 31; hes 14; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 44:
LLECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: 3.5 inch 1.44Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                               Application US/08484138
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30 Rockefeller Plaza
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Weil, Roger
                   Ren, Wu-Yun
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                                  Watanabe, Kyoichi A.
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Roger
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CORRESPONDENCE ADDRESS NUMBER OF SEQUENCES:

INVENTION:

Complementary DNA and Toxins

0

ADDRESSEE: Cooper & Dunham LLP STREET: 1185 Avenue of the Americas

New York New York

U.S.A

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SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,138
FILING DATE: June 7, 1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 44683-Z/JPR
TELECOMMUNICATION INFORMATION:
TELEPAN: 212-977-9550
TELEPAN: 212-977-9550
TELEPAN: 212-978-9550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
            FILING DATE: August 15
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: 08
                                                                                                                                 SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
TUMBER OF SEQUENCES:
                                                                                                                                                           COMPUTER: IBM COM
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                     CITY: Los Angeles
STATE: California
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STRANDEDNESS: dou
STRANDEDNESS: dou
TOPOLOGY: linear
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OPERATING SYSTEM:
                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                                                       Suite 4700
                                                                                                                                                                                                                                                                                                                                      в: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                 IBM Compatible
                                                                                                August 15, 1994
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May 18,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.5%; Score 13.4; DB 1; 93.3%; Pred. No. 31; ative 0; Mismatches 1;
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DISEASES OR CONDITIONS
RELATED TO LEVELS OF
                                                                                                                                                                    IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                          830
                08/245,466
                                                                                                                US/08/291,932A
                                including application described below:
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US-08-442-461D-16
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                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                       TELEFAX: (415) 576-03
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Floppy disk
COMPUTER: POSTM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/442,461D
APPLICATION NUMBER: US/08/442,461D
FILING DATE: 17-MAY-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS; both
                                                                                         TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
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                                                                                                                                                                                                                                     NAME: Liebeschuetz, Joe
REGISTRATION NUMBER: 37,505
REFERRENCE, DOCKET NUMBER: 02
LECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 20
LECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DDRESSEE: Townsend and Townsend and Crew LLP IREET: Two Embarcadero Center, Eighth Floor Sign San Francisco
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 Conservative
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Martin, Shelley S
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                                                                                                             linear
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0.5%; Score 13.4; I
86.7%; Pred. No. 31;
ative 1; Mismatches
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                                 Length 15;
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Best Local Similarity 93.3%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
LENGTH: 15 base pairs
TYPE: NUCLEIC ACID
TYPE: NUCLEIC ACID
                                                                                                                                                TITLE OF INVENTION: Complementary DNA and Toxins NUMBER OF SEQUENCES: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Murashige, Kate H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-327-7250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FO-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch 1.44Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
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                                                                                                                                     ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: POFILING DATE: 19910524
                                                    COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
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TLE OF INVENTION: CROSSLINKING AGENTS WHICH BIND TO THE MAJOR GROOVE OF
TLE OF INVENTION: DUPLEX DNA
                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
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7: 545 Middlefield Road, Suite 200
Menlo Park
California
                                                                                                                                                                                                                                                                                                                                          AAGAGGAGGAGGAGG 15
                                                                    New York
                                                                                                   E: Cooper & Dunham LLP
1185 Avenue of the Americas
                                                                                                                                                                                      Watanabe, Kyoichi A.
Ren, Wu-Yun
Weil, Roger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                415-327-2951
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Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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5217867-3
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Best Local S
Matches 14
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                                                                                                                                                                                                                                     Patent No.
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RESULT 82
US-08-152-019A-13/c
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TITLE OF INVENTION: RECEPTORS THE:
CHARACTERIZATION, PREPARATION AND USE
NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCT-US95-06379-19
                                                                                                                                                                                                                              Sequence 13, Application US/08152019A Patent No. 5565331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 212-391-0526 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Rela
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PC:
FILING DATE: May 13, 1:
                                                     NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 441
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-278-0400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/07/278,614 FILING DATE: 30-NOV-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: double
                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                   1469 GGACCACCATGGGGC 1483
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                                                                                                                                                                                                             INFORMA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5217867
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SSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
C: 4 Embarcadero Center, Suite 3400
San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGAGGAGGAGGAGG 15
                                                                                          INVENTION:
                                                                                                                      Serafini, Tito
Kennedy, Timothy
Placzek, Marysia
Jessell, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EVANS, RONALD M.; HOLLENBERG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15 base pairs
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                                                                                                       Dodd, Jane
                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                    93.3%;
                                                                                       NEURAL AXON OUTGROWTH MODULATORS
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Pred. No. 3
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TELEX: 910 277299 FHT UR
INFORMATION FOR SEQ ID NO: 13:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/152,019A
FILING DATE: 12-NOV-1993
CLASSIFICATION: 514
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: 1i
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                                                                                                         CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                           NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 480124.403C1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
ZIP: 941
                                                                                                                                                                                                      ZIP: 98104-7092
MPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
COPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                      CITY: Seattle
STATE: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Osman, Richard Aron REGISTRATION NUMBER: 36,6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER:
                                                                                                                                        APPLICATION NUMBER: US/08/954,210 FILING DATE: 20-OCT-1997
                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                        STREET:
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          TELEPHONE: (206)
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AL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                       ESPONDENCE
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94111-4187
R READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                        6300 Columbia Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Barber, Jack R. Welch, Peter J. Tritz, Richard
                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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SEED and BERRY LLP
                               (206)
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ION: HEPATITIS C VIRUS RIBOZYMES
NCES: 73
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            6) 622-4900
682-6031
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Pred. No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Hepatitis C Virus
US-09-431-419A-59
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SEQ ID NO 59
LENGTH: 16
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Best Local Similarity 87.9
Matches 14; Conservative
                                                                                            SOFTWARE: Pa
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APPLICANT: PHILLIPS, M.
APPLICANT: ZHANG, YUAN
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                       Patent No. 64893
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OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE -09-614-034-110
                                                                                                                                          PRIOR APPLICATION NUMBER: PCT/US99/21007 PRIOR FILING DATE: 1999-09-14
                                                                                                                                                                                                                                                                                                                                                                    09-614-034-110
                                                                                                                           NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/431,419A
CURRENT FILING DATE: 1999-11-01
                                                                                                                                                                             PRIOR FILING DATE: 1998-09-14
                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/152,717
                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/614,034
CURRENT FILING DATE: 2000-07-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tent No
                                              ORGANISM: UNKNOWN
                                                                                                                                                                                                                                                        ITLE OF INVENTION: ANTISENSE COMPOSITIONS TARGETED TO ITLE OF INVENTION: METHODS OF USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                          E REFERENCE: 4300.013900
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                     16 TCACCTTTGACAGACT 1
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                                                                                                          PatentIn version 3.0
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Pred. No. 4
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Pred. No. 4
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RESULT 87
US-09-705-267A-173/c
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                                                             Sequence 173, Application US/09705267A Patent No. 6551826 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6686188
SEQ ID NO 8648
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                               LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
-09-866-108A-8648
                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Aeomica Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 15755
APPLICANT: Hong Zhang
APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF RAIDD EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/866,108A
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FILE REFERENCE: AEOMICA-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NT FILING DATE: 2001-05-25
APPLICATION NUMBER: US 60/207,456
FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US01/00666
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US01/00669
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                                                                                                                                                                                                                  1652 CCAGCTGCAGAGGCAG 1667
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                                                                                                                                                                                16 CCAGCTGCAGCTGCAG 1
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LING DATE: 2001-01-30
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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ilarity 87.5%;
Conservative
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87.5%;
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Pred. No. 48;
0; Mismatches
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Pred. No. 5
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                                                               US-08-793-660B-22
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Best Local S
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LENGTH: 20

TYPE: DNA

ORGANISM: Artificial Sequence
Query Match 0.9
Best Local Similarity 92.9
Matches 13; Conservative
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CURRENT APPLICATION NUMBER: US/09/705,267A
CURRENT FILING DATE: 2000-11-01
NUMBER OF SEQ ID NOS: 177
                                                                                                                                    TELEFAX: 617-523-6440
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 47
TELECOMMUNICATION INFORMATION:
                                                                           TOPOLOGY: 1 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MULLER, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                          FILING DATE: 26-AUG-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                              FILING DATE: 29-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 94/1
                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 09-SEP-1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U
ZIP: 02109
                                                                                                        LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                NAME: Lowen, Cara Z.
REGISTRATION NUMBER: 38,227
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 96/0:
FILING DATE: 07-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                    CELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: DIKE, BRONSTEIN, ROBERTS & CUSHMAN, LLP 130 Water Street
                                                                         linear
PE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                       617-523-3400
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CELL CYCLE REGULATED REPRESSOR
AND DNA ELEMENT
                0.5%;
92.9%;
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                Score 12.4;
Pred. No. 38
                                                                                                                                                                                                                                     47211
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Pred. No. 83;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Version
                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           #1.30
                              Length 14;
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Mismatches

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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: F
SEQ ID NO 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence:probe sequence -10-032-307-64
                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/054,830 PRIOR FILING DATE: 1998-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 17682A-003630US
CURRENT APPLICATION NUMBER: US/10/032,307
CURRENT FILING DATE: 2001-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 09/431,385
PRIOR FILING DATE: 1999-11-01
PRIOR APPLICATION NUMBER: US 60/186,046
PRIOR FILING DATE: 2000-03-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/054,832
PRIOR FILING DATE: 1998-04-03
                                                                                                                                                                                                                                          quence 247, Application US/08050073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 09/640,953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPLICANT: Vermeulen, Nicolaas M.J.
PPLICANT: Epoch Biosciences, Inc.
TILE OF INVENTION: T-m Leveling Methods
                                                                                                                               APPLICANT:
                                                                                                              APPLICANT:
                                                                                                                                                                     APPLICANT:
                                                  PPLICANT: Scharf, Stephen J.
IITLE OF INVENTION: Methods and Reagents for HLA DRBeta DNA
IITLE OF INVENTION: Typing
IMMBER OF SEQUENCES: 315
                                                                                                                                                                                       PPLICANT:
                                                                                                                                                                                                       PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CANT: Dempcy, Robert ...
CANT: Gall, Alexander A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        no. 6683
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APPLICATION NUMBER: US 09/796,988
                                                                                                                                                                                                                                                                                                                                                                                               1520 CGGCAACGTGCTGG 1533
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APPLICATION NUMBER: US 09/724,959
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Afonina, Irina A.
Singer, Michael J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/10032307
                                                                                                                             Begovich, Ann B.
Bugawan, Teodorica L.
Erlich, Henry A.
Griffith, Robert L.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
Hoffmann-La Roche Inc
0 Kingsland Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001-02-28
                                                                                                                                                                                                       Raymond J.
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Pred. No. 38;
0; Mismatches
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RESULT 91
US-08-166-664-9/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (510) 814-2977
INFORMATION FOR SEQ ID NO: 24
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vatent No. 5646020
GENERAL INFORMATION:
APPLICANT: James
                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARB: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM FOCOMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                            FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Petry, Douglas A.
REGISTRATION NUMBER: 35,321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: J. Anthony Mamone
TITLE OF INVENTION: HAMMERHEAD RIBOZYMES FOR
TITLE OF INVENTION: PREFERRED TARGETS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 87
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                             COUNTRY: U
ZIP: 90017
                                                                                                                                                                                                                                                                                                                                                                    STREET: bil ...
CITY: Los Angeles
C-lifornia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 435
                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
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                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                      DDRESSEE:
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                                                                                                                                                                                                                                                                                                                                              California
: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08166664
                                                                                                                                                                                                                                                                                                                                                                                                   E: Lyon & Lyon
611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               James A. McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (510)
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Pred. No. 47;
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Gaps

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RESULT 92
US-08-291-932A-201/c
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US-08-291-932A-201
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Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 201, Application US/08291932A
Patent No. 5658780
                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                       TELEFAX: (213) 955-044

TELEX: 67-3510

INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AMEAU.
APPLICANT: Successive APPLICANT: Draper, Kenn McSwiggen, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA;
                                                                                                                         REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELLEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                      APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REPLICANT: McSwiggen, James
IITLE OF INVENTION: RIBOZYME TREATMENT OF
IITLE OF INVENTION: DISEASES OR CONDITIONS
IITLE OF INVENTION: RELATED TO LEVELS OF
IITLE OF INVENTION: NF-KB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                   NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                                                          ENGTH:
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                                                          15 base pairs
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Draper, Kenneth G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (213) 955-0440
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               linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                 UMBER: US/08/291,932A
August 15, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single
                                                                                                                                                                                                                                                                                                      including application described below:
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                                                                                            201:
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                                                                                                                                                                         208/157
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Pred. No. 4
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                                                                                          ; TOPOLOGY:
US-08-292-620A-324
                                             Query Match
Best Local
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                               Matches
                                                                                                                                                   TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA: including PRIOR APPLICATION DATA: described APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME
TITLE OF INVENTION: DIEBASES
TITLE OF INVENTION: RELATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                FILING DATE: December 7, 1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSES: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: August CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                             Match 0.5%;
Local Similarity 92.9%;
                                                                                                                          STRANDEDNESS:
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1122 GTCCTCCAAGACCT 1135
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1 Similarity 92.9%;
13; Conservative
                                                                                                                                      nucleic acid
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Dan T. Stinchcomb
James McSwiggen
                               Conservative
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                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                    August 17,
                                                                                                                          single
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DISEASES OR CONDITIONS
RELATED TO LEVELS OF
INTRACELLULAR ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             US/08/292,620A
17, 1994
                                                                                                                                                                                                                                                                                                                                                                                             including application described below:
                              Score 12.4; DI
Pred. No. 47;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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Pred. No. 4
                                                                                                                                                                                                                                                                      208/149
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                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
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                                                            Length 15;
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RESULT 95
US-09-081-646-33
; Sequence 33, Application US/09081646
; Patent No. 633152
; GENERAL INFORMATION:
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US-09-071-845-324/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-071-845-324
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Best Local Similarity
Matches 13; Conserv
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LENGTH: 15 harmonian for the formation of the for
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APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME T.
TITLE OF INVENTION: DISEASES OF
TITLE OF INVENTION: RELATED TO
TITLE OF INVENTION: MOLECULE-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,6
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
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: Los Angeles
3: California
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633 West Fifth Street
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James McSwiggen
Sean Sullivan
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DISEASES OR CONDITIONS
RELATED TO LEVELS OF
INTRACELLULAR ADHESION
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92.9%; Pred. No. 4
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                                                                                                                                                                                                                                                                                                                       8
                                                                                                                                                                                                 RESULT 97
US-08-687-551-10
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FILE REFERENCE: 01107.74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 391
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 33
LENGTH: 15
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APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
APPLICANT: Zhou, Wei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.5%;
Best Local Similarity 92.9%;
Matches 13; Conservative
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Best Local Similarity
Matches 13; Conserv
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TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107.74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER FILING DATE: 1997-05-21
EARLIER FILING DATE: 1997-05-21
MUMBER OF SEQ ID NOS: 871
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                                                                                                                                           GENERAL INFORMATION:
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APPLICANT: EMILE, CArole
APPLICANT: HELENE, Claude
APPLICANT: SPENLEHAUER, Gilles
APPLICANT: SPENLEHAUER, Gilles
TITILE OF INVENTION: NUCLEIC ACID-CONTAINING COMPOSITION,
TITILE OF INVENTION: PREPARATION AND USE
NUMBER OF SEQUENCES: 16
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                                                                                                                     BAZILE, Didier
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                    0.5%; Score 12.4;
92.9%; Pred. No. 4
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                         Length 15;
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Best Local Similarity
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APPLICATION UNMBER: FR 94/013
FILING DATE: 08-FEB-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: W0 PCT/FR
FILING DATE: 27-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Smith Ph.D., Julie K.
REGISTRATION NUMBER: 38,619
REFERENCE/DOCKET NUMBER: ST94
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d:
COMPUTER: IBM PC comp
COMPUTER: PC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (610)454-380
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DESCRIPTION: /desc = "oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/687,551
               ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                         STREET: 633 ....
STREET: Suite 4700
STREET: Los Angeles
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                  APPLICANT: McSwiggen, James
INITLE OF INVENTION: RIBOZYME TREATMENT OF
INITLE OF INVENTION: DISEASES OR CONDITIONS
INTLE OF INVENTION: RELATED TO LEVELS OF
INTLE OF INVENTION: NF-KB
                                                                                                                                                                                                                                                                                                                                                                                     nt No.
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CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 500 Arcola Rd. 3C43
CITY: Collegeville
                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                 ADDRESSEE:
COMPUTER:
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56587,
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                                                                                                                                                              3: Lyon & Lyon
633 West Fifth Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                              Stinchcomb, Dan T.
Draper, Kenneth G.
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100.0%; Pred. No.
                                                                                                                                                                                                                       830
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US-08-291-932A-249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Draper, Kenn APPLICANT: Draper, Kenn APPLICANT: McSwiggen, Critical OF INVENTION: RIB
APPLICATION NUMBER: US FILING DATE: August 15.
CLASSIFICATION: 514
PRIOR APPLICATION DATA: PRIOR APPLICATION DATA: PRIOR APPLICATION DATA: 08.
APPLICATION NUMBER: 08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA: including
PRIOR APPLICATION DATA: describe;
APPLICATION NUMBER: 08/245,466
                                                                                                                                                                                                                                   SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 633 West F11
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ITLE OF INVENTION: RIBOZYME TREATMENT OF ITLE OF INVENTION: DISEASES OR CONDITIONS ITLE OF INVENTION: RELATED TO LEVELS OF TRADED OF CONTROL OF TREATMENT OF THE OF TRADED OF TREATMENT OF TREATMENT OF TRADED OF TRADED
                                                                                                                                                                                                                     COMPUTER: LDW. COMPUTER: LDW. IBW. L..
OPERATING SYSTEM: IBW. L..
OPERATING SYSTEM: IBW. L..
OPERATING DATA:
OPERATING DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: May 18, 1994
APPLICATION NUMBER: 07/987,132
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633 West Fifth Street
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Draper, Kenneth G.
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N: 514
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   : including application
: described below:
   08/245,466
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FILING DATE: May 18
APPLICATION NUMBER:

December 7, 1992

07/987,132

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RESULT 100
US-08-585-684B-2053/c
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Best Local Similarity 83.3%;
          TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENCTH: 15 base pairs
TYPE: nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 5877021 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 489-160
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
                                                                                                        NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
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TITLE OF INVENTION: I
TITLE OF INVENTION: F
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REFERENCE/DOCKET NUMBER: 200
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                        APPLICATION NUMBER: 60/000,951 FILING DATE: July 7, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/585,684B FILING DATE: January 16, 1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
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STRANDEDNESS:
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Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90071
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633 West Fifth Street
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INDUCTION OF GRAFT TOLERANCE
AND REVERSAL OF IMMUNE RESPONSES
2751
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Pred. No. 55;
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RESULT 101
US-08-271-882B-30
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TOPOLOGY:
US-08-271-882B-30
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                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: . . SEQUENCE CHARACTERISTICS: LEUGTH: 15
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                                                                 Matches
                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Glen A. Evans
APPLICANT: Ronald G. Sosnowski
TITLE OF INVENTION: SELF-ADDRESSABLE
TITLE OF INVENTION: MICROELECTRONIC SY.
TITLE OF INVENTION: DEVICES FOR
TITLE OF INVENTION: MOLECULAR BIOLOGIC.
TITLE OF INVENTION: AND DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/271,882B
FILLING DATE: July 7, 1994
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                   NAME: Murphy, David B.
REGISTRATION NUMBER: 31,125
REFERENCE/DOCKT NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/146,504
FILING DATE: No. 6017696ember 1, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Mulphy, David B.
                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
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                                                                                                                                                 STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                                                                                   TYPE: nucleic
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[ATE: California
                                1791 GCGGCAGCCACA 1802
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3 GCGGCAGCCACA 14
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633 West Fifth Street
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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                                                                                 0.5%;
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                                                                                   Score 12;
pred. No.
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55;
                                                                                       DB 1;
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                                                                                                 Length 15
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RESULT 103
US-09-275-850-21
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APPLICANT: Gold, Larry
APPLICANT: Shtatland, Timur
APPLICANT: Shtatland, Timur
APPLICANT: Javornik, Brenda
TITLE OF INVENTION: Truncation SELEX Method
FILE REFERENCE: NEX 79
CURRENT APPLICATION NUMBER: US/09/275,850A
CURRENT FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 351
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-038-073-2053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.5%; Score 12; Best*Local Similarity 100.0%; Pred. No. Matches 12; Conservative 0; Mismatc
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 15 base --
                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21, Appearance 21, Appearan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FASTSEQ Version 1.
CURRENT APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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10. 6194150
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633 West Fifth Street
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                                                                                                                                                                                                  Truncation SELEX Method
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 105
US-08-486-421-51/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Hong Zhang
APPLICANT: Susan M. Freier
APPLICANT: SUSAN M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF RAIDD EXPRESSION
FILE REFERENCE: RTS-0211
CURRENT APPLICATION NUMBER: US/09/705,267A
CURRENT FILING DATE: 2000-11-01
NUMBER OF SEG ID NOS: 177
SEG ID NO 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 86.,
"Arches 13; Conservative
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LENGTH: 15
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Sequence 51, App-
Sequence 51, App-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 20
                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: LEM PC compatible
COMPUTER: PEC COMPATIBLE
COMPUTER: PC COMPATIBLE
COPERATING SYSTEM: PC CDS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/486,421
APPLICATION NUMBER: US/08/486,421
                                                                                                                                                                                                                                                                                                                      APPLICANT: Johnson, Edward M.
APPLICANT: Bergemann, Andrew D.
TITLE OF INVENTION: CLONING AND EXPRESSION OF PUR PROTEIN
NUMBER OF SEQUENCES: 51
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/470,911
FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                          STREET: 1155 Av
CITY: New York
STATE: New Yorl
                                                                      FILING DATE:
                                                                                                                                                                                                         COUNTRY: U.S.A.
ZIP: 10036-2711
                                                                                                                                                                                                                                                                                              DDRESSEE:
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1155 Avenue of the Americas
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                                                                  UMBER: US/08/486,421
07-JUN-1995
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; Pred. No. 55;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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Pred. No. 1.1e+02;
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; MOLECULE TYPE: DNA
US-08-486-421-51
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Best Local Similarity 69.0
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equence 51, April
Mo. 5756684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-470-911-51/c
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 692:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE Floppy disk
COMPUTER: Floppy disk
COMPUTER: FR PC compatible
COMPUTER: PATENT PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
110 / 10 / 10 / 10 0 11
                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Johnson, Edward M.
APPLICANT: Bergemann, Andrew D.
TITLE OF INVENTION: CLONING AND EXPRESSION OF PUR PROTEIN
NUMBER OF SEQUENCES: 51
TOPOLOGY: 1:
                                                                                                                                                                      NAME: COPUZZÍ, LAURA À.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 69.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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                               STRANDEDNESS:
                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/470,911 FILING DATE: 06-JUN-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RRESPONDENCE ADDRESS:
                                                                                                                   TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
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1155 Avenue of the Americas
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RESULT 107
US-08-486-809-51/c
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US-09-156-979-46
; Sequence 46, A
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Patent No. 5869622
                                             GENERAL INFORMATION:
APPLICANT: COWSET, Lex M.
APPLICANT: COWSET, Lex M.
TITLE OF INVENTION: ANTISENSE MODULATION (
FILE REFERENCE: RTS-0013
CURRENT FILING DATE: 1998-09-18
NUMBER OF SEQ ID NOS: 47
SEQ ID NO 46
SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local :
                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (212) 869-974
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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ORGANISM: Artificial Sequence
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MIDUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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APPLICANT: Bergemann, Andrew D.
APPLICANT: TUVENTION: CLONING AND EXPRESSION OF PUR PROTEIN
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PRIOR APPLICATION NUMBER: US 08/470,911
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APPLICATION NUMBER: US
FILING DATE: 07-JUN-19
                  TYPE: DNA
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REFERENCE/DOCKET NUMBER: 69
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
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1155 Avenue of the Americas
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Pred. No. 1
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; FEATURE: ; OTHER INFORMATION: Antisense Oligonucleotide US-09-156-979-46
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US-09-387-341-107
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EARLIER FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 233
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 107
LENGTH: 18
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Best Local Similarity 77.8
Matches 14; Conservative
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Best Local Similarity
Matches 14; Conserv
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART.
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-05-26
PRIOR PELING DATE: 2000-05-26
PRIOR PELING DATE: 2000-05-26
PRIOR PILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-05-26
PRIOR PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PILING DATE: 2000-00-04
PRIOR PILING DATE: 2000-00-04
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atent No. 6686188
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EARLIER FILING DATE: 1998-09-18
EARLIER APPLICATION NUMBER: 09/156,979
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CURRENT APPLICATION NUMBER: US/09/387,341
CURRENT FILING DATE: 1999-08-31
                                                                                                                                                                                                                                                                                                                         APPLICANT: GU, Yizhong
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PPLICANT: Cowsert, Lex M.
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APPLICATION NUMBER: 09/156,807
FILING DATE: 1998-09-18
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PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
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Pred. No. 98;
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Pred. No. 98;
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; SOFTWARE: Acomica Sequence Listing Engine
patent NO. 6686188
SEQ ID NO 7854
LENGTH: 17
TYPE: DNA
CORGANISM: Homo sapiens
US-09-866-108A-7854
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                                                                                       Matches 13; Conservative
                                                                                                           Query Match 0.4%;
Best Local Similarity 81.2%;
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PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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(S-10-308-503-110)
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$-09-864-785-1560
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$-09-780-533A-2363
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                 ; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:
; OTHER INFORMATION: oligonucleotide
US-10-275-071-18
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US-10-275-071-18/c
                                                                           ; TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
US-09-993-731-36
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CURRENT APPLICATION NUMBER: US/10/275,071
CURRENT FILING DATE: 2003.04-07
PRIOR APPLICATION NUMBER: 09/580,923
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 08/860,038
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 1997-06-09
PRIOR APPLICATION NUMBER: PCT/FR95/01468
PRIOR FILING DATE: 1995-11-08
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
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                                     Query Match
Best Local S
Matches 20
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Best Local S
Matches 21
                                                                                                                          APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 36
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18, Application US/10275071
Publication No. US20030186286A1
GENERAL INFORMATION:
APPLICANT: Crouzet, Joel
APPLICANT: Scherman, Daniel
APPLICANT: Scherman, Daniel
                                                                                                                                                                                                                 Sequence 36, Application US/09993731 Publication No. US20030105040A1 GENERAL INFORMATION:
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SEQ ID NO 18
LENGTH: 25
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APPLICANT: Blanche, Francis
TITLE OF INVENTION: PURIFICATION OF A TRIPLE HELIX FORMATION WITH
TITLE OF INVENTION: IMMOBILIZED OLIGONUCLEOTIDE
                                                                                                                                                                                                                                                                                                       1771 AGGAGGAGGAGGCGAGGC 1792
                   840 CTGAATGAGATGAGGACCCG 859
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  CTGAATGAGATGAGGACCCG
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Wils, Pierre
                                     0.8%;
ilarity 100.0%;
Conservative
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                                                 Score 20;
Pred. No.
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Pred. No. 6.
                                        Mismatches
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RESULT 3 US-09-993-731-37/c

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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 37
LENGTH: 20
TYPE: DIA
ORGANISM: Artificial Sequence
FEATURE:
FRATURE:
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US-09-993-731-38
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                                        NUMBER OF SEQ ID NOS: 89
SEQ ID NO 39
LENGTH: 20
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Publication No. US20030105040A1
GENERAL INFORMATION:
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Publication No. US20030105040A1
                                                                                                APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 38
LENGTH: 20
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENERAL INFORMATION:
                                                                                                                                                                         APPLICANT: Brett
APPLICANT: Andre
TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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ORGANISM: Artificial Sequence
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20; Conservative 0;
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; Pred. No. 4.1;
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Pred. No.
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                                                                                                                  ; OTHER INFORMATION: US-09-993-731-41
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LENGTH: 20
TYPE: DNA
                                                      Query Match
Best Local S
Matches 20
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Best Local S
Matches 20
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Publication No.
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TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                    APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                               APPLICANT: Brett P. Monia APPLICANT: Andrew T. Wat
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APPLICANT: Andrew
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                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                              FEATURE:
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                                                                                                                                                                                         ENGTH: 20
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                           960 CTTTACGAGGACCTATTCCG 979
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Pred. No.
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RESULT 8
US-09-993-731-42/c
; Sequence 42, Application US/09993731

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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-993-731-42
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US-09-993-731-44/c
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US-09-993-731-43/c
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Best Local S
Matches 20
                                                                  APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 44
LENGTH: 20
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 42
LENGTH: 20
TYPE-CARP
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Publication No. US20030105040A1
GENERAL INFORMATION:
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SEQ ID NO 43
LENGTH: 20
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Best Local Similarity
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GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-11-13
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
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                                                        TYPE: DNA
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                 FEATURE:
                                   ORGANISM: Artificial Sequence
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INFORMATION: Antisense Oligonucleotide
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RESULT 13
US-09-93-731-47/c
Sequence 47, Application US/09993731
; Publication No. US20030105040A1
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                                                                                                                                                                                                                                ; OTHER INFORMATION: Antisense Oligonucleotide US-09-993-731-46
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 46
LENGTH: 20
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SEQ ID NO 45
LENGTH: 20
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Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
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                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                  Local Similarity
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100.0%; Pred. No.
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100.0%; Pred. No.
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Best Local Similarity
Matches 20; Conservi
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US-09-993-731-49/c
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SEQ ID NO 47
LENGTH: 20
TYPE
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 48
LENGTH: 20
TYPET - 120
                                                                                                           Sequence 49, Application US/09993731

Publication No. US30030105040A1

GENERAL INFORMATION:

APPLICANT: Erett P. Monia

APPLICANT: Andrew T. Watt

TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION

FILE REFERENCE: RTS-0302

CURRENT APPLICATION NUMBER: US/09/993,731

CURRENT FILING DATE: 2001-11-13

NUMBER OF SEQ ID NOS: 89
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Best Local Similarity
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Sequence 48, Application US/09993731
Publication No. US20030105040A1
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TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION FILE REFERENCE: RTS-0302
OTHER INFORMATION: Antisense Oligonucleotide 09-993-731-49
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APPLICANT: Andrew T. Wat
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                                                               EQ ID NO 49
LENGTH: 20
TYPE: DNA
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ORGANISM: Artificial
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                             FEATURE:
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Pred. No.
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US-09-993-731-52/c
                RESULT 18
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 51
LENGTH: 20
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LENGTH: 20
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Best Local
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TITLE OF INVENTION: ANVISENSE MODULATION OF INHIBITOR-KAPPA
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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ORGANISM: Artificial Sequence
FEATURE:
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100.0%; Pred. No. 4.1;
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Pred. No.
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Sequence 52, Application US/09993731 Publication No. US20030105040A1

GENERAL INFORMATION

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                                                                                                                   GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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SEQ ID NO 52
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
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Best Local Similarity
                                                                                   SEQ ID NO 54
LENGTH: 20
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Publication No.
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OTHER INFORMATION: Antisense Oligonucleotide -09-993-731-54
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CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O ID NO 53
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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                                    FEATURE:
                                                  ORGANISM: Artificial Sequence
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20; Conserv
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lo. US20030105040A1
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100.0%; Pred. No. 4.1;
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L00.0%; Pred. No.
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RESULT 23
US-09-993-731-57/c
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Publication No. US20030105040A1

GENERAL INFORMATION:

APPLICANT: Brett P. Monia

APPLICANT: Andrew T. Watt

TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION

FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                            FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 56
LENGTH: 20
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Best Local Similarity
Matches 20; Conserv
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Best Local S
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LENGTH: 20
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Best Local Similarity
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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TYPE: DNA
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ITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
                                                                                          1258 AGCAACAGCTGGAAGAGGCT 1277
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100.0%; Pred. No. 4.
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Sequence 57, Application US/09993731 Publication No. US20030105040A1 GENERAL INFORMATION:

APPLICANT: Brett P.

Monia

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US-09-993-731-59/c
Sequence 59, Application US/09993731
Publication No. US20030105040A1
GENERAL INFORMATION:
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Best Local Similarity 100.
20; Conservative
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Publication No. US20030105040A1
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
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CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 57
LENGTH: 20
Query Match
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                         FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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                                                                                                                                                                                                                                APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
                                                                                                       EQ ID NO 59
LENGTH: 20
TYPE: DNA
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
                                                                      ORGANISM: Artificial Sequence FEATURE:
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ORGANISM: Artificial
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                                                   OTHER INFORMATION: Antisense Oligonucleotide
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 Score 20; DB 1;
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Length 20;
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RESULT 28
US-09-993-731-62/c
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-993-731-60
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Best Local S
Matches 20
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Best Local S
Matches 20
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
TITLE PRIFERENCE: RTS-0302
CURRENT FILLYS DATE: 2001-11-13
CURRENT FILLYS DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 60
SEQ ID NO 60
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Publication No. US20030105040A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                           SEQ ID NO 61
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                                                                   APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                              FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                            TYPE: DNA
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4.1;
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; Sequence 62, Applicat:
; Publication No. US200;
; GENERAL INFORMATION:
; APPLICANT: Breett P. !
; APPLICANT: Andrew T

Application US/09993731 to. US20030105040A1

Brett P. Monia Andrew T. Watt

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; TYPE: DNA; ORGANISM: Artificial Sequence; PEATURE: TYPORMATION: Antisense Oligonucleotide US-09-993-731-62
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US-09-993-731-64/c
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US-09-993-731-63/c
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                                                FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-993-731-64
                                                                                                                                                    APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0302
CURRENT FILING DATE: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
1500 NO 640
                                                                                                                                                                                                                                                                                                  Sequence 64, Application US/09993731
Publication No. US20030105040A1
GENERAL INFORMATION:
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Best Local Similarity 100.0%;
Matches 20; Conservative (
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LENGTH: 20
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Query Match
Best Local Similarity
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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APPLICANT: Andrew T. Wart
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION FILE REFERENCE: RTS-0302
                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                            ENGTH:
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 Score 20;
Pred. No.
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 DB 1;
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                                              US-09-993-731-67/c
, Sequence 67, Application US/09993731
, Publication No. US20030105040A1
, GENERAL INFORMATION:
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                                                                                                              RESULT 33
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Best Local Similarity
Matches 20; Conserv
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LENGTH: 20
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Best Local
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA
FILE REFERENCE: RTS-0302
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0302
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Brett P. Monia APPLICANT: Andrew T. Wat
                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
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No. US20030105040A1
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4.1;
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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TYPE: DNA
ORGANISM: Artificial S
FEATURE:
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SEQ ID NO 69

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Publication No. US20030105040A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 20; Conserv
Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Fred. No. 4.1; Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                               APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Wastt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
FULL REFERENCE: RTS-0302
CURRENT FAPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                             OTHER INFORMATION: -09-993-731-69
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CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                                               APPLICANT: Brett APPLICANT: Andre
                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                    FEATURE:
                                                                                                                                                                            ENGTH: 20
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                                                                                                Antisense Oligonucleotide
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100.0%; Pred. No. 4.1;
ative 0; Mismatches
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FILECANT: ADDREW T. WALE
FILE OF INVENTION: ANTISENSE MODULATION OF INE
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
FIRMOTE:
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                                                                                                            US-09-993-731-72/c
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US-09-993-731-71/c
                                                                                                                            RESULT 38
Sequence 72, Application US/09993731
Publication No. US20030105040A1
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 71, Application US/09993731
Publication No. US20030105040A1
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 70
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APPLICANT: Brett P.
APPLICANT: Andrew
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                              LENGTH: 20
TYPE: DNA
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                                                                              Application US/09993731
No. US20030105040A1
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Pred. No.
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4.1;
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US-09-993-731-73/c
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                                                               ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-993-731-74
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Best Local S
Matches 20
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 74
SEQ ID NO 74
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Publication No. US20030105040A1
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
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Best Local
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Best Local S
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 73
                                                                                                                                                                                                                             APPLICANT: ANDREW T. WALL
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
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               100.0%;
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100.0%; Pred. No. 4.1;
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               DB 1;
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TITLE OF INVENTION: ANTISENSE MODULATION OF IN:
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 76
LENGTON
                                              US-09-993-731-77/c
; Sequence 77, Application US/09993731
; Publication No. US20030105040A1
; GENERAL INFORMATION:
; APPLICANT: Brett P. Monia
; APPLICANT: Andrew T. Watt
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US-09-993-731-76/c
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Best Local Similarity

Matches 20; Conserv
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Publication No. US2
GENERAL INFORMATION
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TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
         APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
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CURRENT APPLICATION NUMBER: US/09/993,731
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100.0%; Pred. No. 4.1;
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Pred. No.
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CURRENT FILING DATE:
NUMBER OF SEQ ID NOS:
SEQ ID NO 77

ENGTH: 20

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TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION FILE REFERENCE: RTS-0302 CURRENT APPLICATION NUMBER: US/09/993,731; CURRENT FILLING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
LENGTH: 20
TYPE: DN7
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                                                                                       ; OTHER INFORMATION: Antisense Oligonucleotide US-09-993-731-79
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                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 79
LENGTH: 20
                                                                                                                                                                                                                                                                  Sequence 79, Application US/09993731
Publication No. US20030105040A1
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
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Best Local Similarity
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                              Matches
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                                                                                                                                                                                                                                       TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION FILE REFERENCE: RTS-0302
                                                                                                                              ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                                    TYPE: DNA
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   1662 AGGCAGGTCTTGCAGCATCT 1681
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-993-731-80
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US-09-993-731-82/c
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; Sequence 81, Application US/09993731;
; Publication No. US20030105040A1
; GENERAL INFORMATION:
; APPLICANT: Brett P. Monia
; APPLICANT: Andrew T. Watt
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US-09-993-731-80/c
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                                                                                                     Sequence 82, Application US/09993731
Publication No. US20030105040A1
GENERAL INFORMATION:
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
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Best Local Similarity
Matches 20; Conserv
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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Publication No. US20030105040A1
GENERAL INFORMATION:
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TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER.OF SEQ ID NOS: 89
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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                                                                        APPLICANT: Brett P. APPLICANT: Andrew
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 83
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Best Local Similarity
Matches 20; Conserv
                                                  Query Match
Best Local Similarity
Matches 20; Conserv
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LENGTH: 20
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Publication No.
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SEQ ID NO 82
LENGTH: 20
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
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APPLICANT: Andro
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TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
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PLICANT: Andrew T. Watt
PLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
E REFERENCE: RTS-0302
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                                                0.8%; Score 20; DB 1; Length 20, 100.0%; Pred. No. 4.1; tive 0; Mismatches 0; Indels
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                                       ; ORGANISM: Homo sapiens
US-10-005-626A-70
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                                                                              SEQ ID NO 70
LENGTH: 23
TYPE: DNA
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Matches 19; Conserv
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Query Match
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Patent No. US20020064771A1
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CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: US. 60/195, 852
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Patentin Ver. 2.1
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NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn vei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 21401-7002
CURRENT APPLICATION NUMBER: US/10/005,626A
CURRENT FILING DATE: 2001-12-03
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PRIOR APPLICATION NUMBER: US 08/682,054
PRIOR FILING DATE: 1996-07-16
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PRIOR FILING DATE: 2001-12-03
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APPLICANT: Hong, Zhi
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ITILE OF INVENTION: Locus Alleles as Haplotypes
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APPLICANT: Michael, Sim
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TITLE OF INVENTION: HCV REPLICASE COMPLEXES
FILE REFERENCE: IN01165
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/551,239 FILING DATE: 1990-07-11
                                                                                                                                                                                                        APPLICATION NUMBER: US 07/398,217
                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 07/949,652
                                                                                                                                                                                                                           FILING DATE: 1989-09-11
                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/405,499
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No. US20030119003A1
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PRIOR FILING DATE: 2002-04-17
NUMBER OF SEQ ID NOS: 423
SOFTWARE: PASTSEQ for Windows Version 4.0
SEQ ID NO 126
                                                                Query Match
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SEQ ID NO 153
LENGTH: 20
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Matches :
                                                                                                                         OTHER INFORMATION: Antisense Oligonucleotide -10-181-846-153
                                                                                                                                                                                                                                                                               FILE REFERENCE: RTSP-0363
CURRENT APPLICATION NUMBER: US/10/181,846
CURRENT FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: PCT/US01/01416
PRIOR FILING DATE: 2001-01-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-181-846-153/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 1551.2001-001
CURRENT APPLICATION NUMBER: US/10/418,182
CURRENT FILING DATE: 2003-04-16
                                                                                                                                                                                                                                                PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: 09/490,692
PRIOR FILING DATE: 2000-01-24
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                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                          CANT: Lex M. Cowsert
OF INVENTION: ANTISENSE MODULATION OF DAXX EXPRESSION
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Pred. No. 1
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Pred. No. 1
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RESULT 55

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SOFTWARE: PatentIn version 3.1
SEQ ID NO 4054
LENGTH: 20
TYPE: DNA
ORGANISM: Candida albicans
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; Sequence 4054, Application US/10032585

; Publication No. US20030180953A1
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NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 9
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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Best Local S
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TITLE OF INVENTION: NUCLEIC ACID DETECTION METHOD
FILE REFERENCE: 109845.191US2; TMRI-020US
CURRENT APPLICATION NUMBER: US/10/388,329
CURRENT FILING DATE: 2003-03-13
CURRENT FILING DATE: 2003-03-13
PRIOR APPLICATION NUMBER: 60/364,230
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APPLICANT: Howard, Bussey
TITLE OF THE TITLE
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                                                                                                                                APPLICANT: Batra, Surinder K.
APPLICANT: Brandt, Randall E.
APPLICANT: Ringel, J'erg
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b. US20020150894A1
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Similarity 94.7%;
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L"hr, Matthias
Varshney, Grish C.
University of Nebraska Board
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94.78;
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RESULT 59
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LENGTH: 18
                                                                Sequence 70, Application US/10199199
Publication No. US20040014047A1
GENERAL INFORMATION:
APPLICANT: Lex M. Cowsert
APPLICANT: Kenneth W. Dobie
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Best Local Similarity
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APPLICANT: University of Nebraska Board of Regents
TITLE OF INVENTION: Specific Mucin Expression as a N
TITLE OF INVENTION: for Pancreatic Cancer
FILE REFERENCE: UNMC 63155
CURRENT APPLICATION NUMBER: US/10/279,454
CURRENT FILING DATE: 2002-10-24
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            APPLICANT: Kenneth M. Dobie TITLE OF INVENTION: ANTISENSE MODULATION OF LIM DOMAIN KINASE 1 EXPRESSION FILE REFERENCE: RTS-0375
CURRENT APPLICATION NUMBER: US/10/199,199
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PRIOR FILING DATE: 2000-12-08
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APPLICANT: Brandt, Randall E.
APPLICANT: Ringel, J"erg
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CURRENT FILING DATE: 2000-12-08
NUMBER OF SEQ ID NOS: 29
SOPTWARE: FastSEQ for Windows Version 3.0
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
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To. US20030134343A1
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Pred. No. 2
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Pred. No. 20
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                                                               ; OTHER INFORMATION: Antisense Oligonucleotide US-10-181-846-155
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APPLICANT: Kenneth W. Dobie
APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF LIM DOMAIN KINASE 1 EXPRESSION
FILE REFERENCE: RTS-0375
CURRENT APPLICATION NUMBER: US/10/199, 199
CURRENT FILING DATE: 2002-07-18
NUMBER OF SEQ ID NOS: 148
SEQ ID NO 135
LENGTH: 20
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Publication No. US20030083297A1
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Publication No.
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SEQ ID NO 70
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                                                                                                                                                             SEQ ID NO 155
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                                Query Match
                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/US01/01416
PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: 09/490,692
PRIOR FILING DATE: 2000-01-24
                                                                                                                                                                                                                                                                                                     APPLICANT: Nicholas M. Dean
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF
                                                                                                                                                                             NUMBER OF SEQ ID NOS: 176
                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/181,846
CURRENT FILING DATE: 2002-07-17
                                                                                                                                                                                                                                                                                          FILE REFERENCE: RTSP-0363
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                                                                                                FEATURE:
                                                                                                               ORGANISM: Artificial Sequence
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Local Similarity tes 16; Conserv
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o. US20040014047A1
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 0.6%; Score 16; DB 1;
100.0%; Pred. No. 33;
ive 0; Mismatches
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Pred. No. 27
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Pred. No. 27;
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Conservative

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RESULT 63
US-09-866-108-8659
; Sequence 8659, Application US/09866108
; Patent No. US20020048800A1
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PELLCATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: DETUNE NUMBER: PETUNG DATE: 2000-09-27
PRIOR APPLICATION NUMBER: DETUNE NUMBER: PETUNG DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engir
SEQ ID NO 929
LENGTH: 17
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Best Local (
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-09-866-108-929
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
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ENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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L, David K.
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Pred. No. 2
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RESULT 64
US-09-780-533A-2359
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-8659
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PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00665
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Best Local S
Matches 16
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Blatt, Larry
                                                                        Sequence 2359, Application US/09780533A Publication No. US20030060611A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 8659
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
APPLICANT: HANZEL, David
APPLICANT: RANK, David R.
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RIOR APPLICATION NUMBER: US 60/266,860 PRIOR FILING DATE: 2001-02-05
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APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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cl Similarity 94.1%;
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EL, David
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Pred. No. 28
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APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pette
TITLE OF INVENTION: Method and Reagent for the Inhibition
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09

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US-09-969-373-3292
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 2359
LENGTH: 17
                                                                   GENERAL INFORMATION:
APPLICANT: Effertz, Roger J.
APPLICANT: Hauge, Brian M.
                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
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Best Local Similarity
                                                                                                                         Sequence 3292, Application US/09969373
Patent No. US20020133852A1
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Publication No.
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TYPE: DNA
ORGANISM: Homo sapiens
-10-060-756A-171
APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)A
CURRENT APPLICATION NUMBER: US/09/969,373
CURRENT FILING DATE: 2001-10-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Aeomica Sequence Listing Engine
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FILING DATE: 2001-10-09
R OF SEQ ID NOS: 4804
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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LING DATE: 2001-01-30
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Similarity 94.1%;
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o. US20030046717A1
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Pred. No. 28;
0; Mismatches
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                  US-10-253-967-39
                                                                                                                                                                                                                                                                                                                                                                     RESULT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-314-405-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/314,405
CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: US/09/713,616
PRIOR FILING DATE: 2000-11-15
NUMBER OF SEQ ID NOS: 46
SOFTWARE: Patentin version 3.0
SEQ ID NO 44
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                      Sequence 39, Application US/10253967 Publication No. US20030165925A1 GENERAL INFORMATION:
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                                                                                         SOFTWARE: PatentIn version 3.1 SEQ ID NO 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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LENGTH: 18
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Best Local Similarity
                                                                                                                                                                                        TITLE OF INVENTION: DIAGNOSTIC PROBE DETECTION SYSTEM FILE REFERENCE: 27978/37504A CURRENT APPLICATION NUMBER: US/10/253,967 CURRENT FILING DATE: 2002-09-24
                                                                                                                              PRIOR APPLICATION NUMBER: US 60/324,421
PRIOR FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: YASUNATI, MATSUZAKA
TITLE OF INVENTION: NOVEL POLYMORPHIC MICROSATELLITE MARKERS IN THE HUMAN MHC CLASS I
FILE REFERENCE: 06501-069001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 09/754,853
PRIOR FILING DATE: 2001-01-05
PRIOR APPLICATION NUMBER: US 09/760,427
PRIOR FILING DATE: 2001-01-13
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APPLICANT: Gen, Tamiya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2001-01-13
PRIOR APPLICATION NUMBER: US 09/855,768
PRIOR FILING DATE: 2001-05-15
                                                                                                                                                                                                                                                                     APPLICANT: SAITO et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 4593
                ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
OTHER INFORMATION: Probe, DQ33
                                                                          ENGTH: 18
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Pred. No. 33;
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Pred. No. 4
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US-10-253-967**-3**9

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RESULT 70
US-10-199-70
(S-10-199-70)
; Sequence 70, Application US/10199199
; Publication No. US20040014047A1
; GENERAL INFORMATION:
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                                      ; OTHER INFORMATION: Antisense Oligonucleotide US-10-199-199-70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence:Synthetic US-10-178-325-107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 16; Conserv
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SEQ ID NO 107
LENGTH: 18
Query Match
                                                                                                                                                 NUMBER OF SEQ ID NOS: 148
SEQ ID NO 70
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                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/199,199
CURRENT FILING DATE: 2002-07-18
                                                                                                                                                                                                                                                                   APPLICANT: Lex M. Cowsert
APPLICANT: Kenneth W. Do
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1998-09-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/178,325
CURRENT FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: US/09/387,341
PRIOR FILING DATE: 1999-08-31
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ablication No.
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                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
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                                                                            FEATURE:
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ITLE OF INVENTION: Expression
ILE REFERENCE: ISPH-0404
                                                                                                                                                                                                                              PPLICANT: Kenneth W. Dobie
TTLE OF INVENTION: ANTISENSE MODULATION OF LIM DOMAIN KINASE 1 EXPRESSION
ILE REFERENCE: RTS-0375
                                                                                                                                   ENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OR APPLICATION NUMBER: 09/156,424
OR FILLING DATE: 1998-09-18
OR FILING DATE: 1998-09-18
OR FILING DATE: 1998-09-18
OR APPLICATION NUMBER: 09/156,807
OR APPLICATION NUMBER: 09/166,807
OR FILING DATE: 1998-09-18
OR APPLICATION NUMBER: 09/161,015
OR FILING DATE: 1998-09-25
OR FILING DATE: 1998-09-25
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88.9%;
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Pred. No. 4
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Pred. No. 44;
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DB 1;
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CURRENT APPLICATION NUMBER: US/10/199,199
CURRENT FILING DATE: 2002-07-18
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PRIO
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TYPE: DNA
ORGANISM: H. sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART
FILE REFERENCE: ADOMICA-7
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Local Similarity 88.9%;
                                                                                                                                                                          APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00662
                                                                                                  APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
                                                                   APPLICATION NUMBER: PCT/US01/00663
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US20020048800A1
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o. US20040014047A1
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, David K.
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
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LENGTH: 17
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
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ORGANISM: Homo sapiens
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PPLICANT: CHEN, Wensheng
PPLICANT: SHANNON, Mark
PTLICANT: SHANNON, Mark
PTLICANT: SHANNON: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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APPLICATION NUMBER: PCT/US01/00666
APPLICATION NUMBER: PCT/US01/00666
                                                   APPLICATION NUMBER: PCT/USO1/00661
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/USO1/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00663
APPLICATION NUMBER: PCT/US01/00663
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
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                                  CATION NUMBER: US 60/266,860
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PENN, Sharron G.
HANZEL, David K.
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2001-02-05
DS: 15752
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Pred. No. 46;
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-930
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                                                                                                     ; ORGANISM: Homo US-09-866-108-2617
Query Match
Best Local Similarity 93.0
Thes 15; Conservative
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PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engi-
SEQ ID NO 2617
LENGTH: 17
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Best Local :
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/236,359
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INVENTION: MYOSIN-LIKE GENE EXPRESSED
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CHEN, Wensheng
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Query Match
Best Local Similarity
Marches 15; Conserva
RESULT 76
US-09-866-108-6391
; Sequence 6391, Application US/09866108
; Patent No. US20020048800A1
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US-09-866-108-2618
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SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 2618
LENGTH: 17
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
PRIOR PELICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PRIOR DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI
APPLICANT: PE
APPLICANT: HA
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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atent No. US20020048800A1
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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PPLICATION NUMBER: PCT/US01/00668
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NN, Sharron G.
NZEL, David K.
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Pred. No. 4
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RESULT 77
US-09-866-108-6392
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; ORGANISM: Homo sapiens
US-09-866-108-6391
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PRIOR FILING DATE: 2000-05-26
PRIOR PPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: CT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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Best Local S
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SOFTWARE: Aeomica Sequence
SEQ ID NO 6391
                                                                                                                                                                                                         Sequence 6392, Application US/09866108 Patent No. US20020048800A1
APPLICANT: PENN, SHARRON G.
APPLICANT: HANKEL, David K.
APPLICANT: RANK David K.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION. MOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE FILE REFERENCE: AEOMICA-7
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: MYOSI
FILE REFERENCE: AEOMICA-7
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CURRENT FILING DATE: 2001-05-25
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FILING DATE: 2001-01-30
ADDITCATE: 2001-01-30
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FILING DATE: 2001-01-30
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                                                                                                                                                 JI, Yonggang
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MN, Sharron G.
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93.8%; Pred. No. 46;
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SOFTWARE: Aeomica Sequence
SEQ ID NO 6392
LENGTH: 17
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                                                              PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/0066
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CURRENT FILING DATE: 2001-05-25
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PPLICANT: HANZEL, David K.
PPLICANT: RANK, David R.
PPLICANT: CHEN, Wensheng
PPLICANT: SHANNON, Mark
ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
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APPLICATION NUMBER: US 60/266,860
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APPLICATION NUMBER: PCT/US01/00666
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1 Similarity 93.8%;
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ICATION NUMBER: PCT/US01/00666
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Pred. No. 46;
0; Mismatches 1;
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SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 8658
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI, Yongga
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CITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
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APPLICATION NUMBER: PCT
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APPLICATION NUMBER: PCT/US01/00668
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PENN, Sharron (
HANZEL, David I
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: 2001-01-30
NUMBER: PCT/US01/00668
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RESULT 80
US-09-825-805-668/c
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US-09-866-108-8660
                                                                                                                                                                      SOFTWARE: PatentIn version 3.0
SEQ ID NO 668
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
                                                             Query Match
Best Local Similarity 93.8
Matches 15; Conservative
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Best Local Similarity 93.8%;
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                                                                                                                                                     -09-825-805-668
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TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleot
FILE REFERENCE: MBHB00-831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 09/578,223
PRIOR APPLICATION NUMBER: 09/476,387
PRIOR FILING DATE: 2000-05-23
PRIOR FILING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/301,511
PRIOR APPLICATION NUMBER: 09/301,511
PRIOR APPLICATION NUMBER: 09/304-28
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR APPLICATION NUMBER: 09/186,675
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PRIOR APPLICATION NUMBER: 60/064,866
PRIOR FILING DATE: 1997-11-05
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                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 1558
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APPLICANT: Beigelman, Leo
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PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
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APPLICATION NUMBER: 60/083,727
1266 CTGGAAGAGGCTGAGG 1281
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Beaudry, Amber
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o. US20030004122A1
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                                                             Score 14.4; DI
Pred. No. 46;
0; Mismatches
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Pred. No. 4
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; ORGANISM: Homo sapiens US-09-818-875-916
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CURRENT APPLICATION NUMBER: US/09/818,875
CURRENT FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR PELING DATE: 2000-03-27
PRIOR PELICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/204,598
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
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US-09-818-875-915
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PRIOR FILLING DATE: 2000-03-27
PRIOR PELICATION NUMBER: US 60/192,179
PRIOR FILLING DATE: 2000-03-27
PRIOR FILLING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILLING DATE: 2000-06-01
PRIOR FILLING DATE: 2000-10-30
                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 4385
SOFTWARE: Friedman macro Napro4
SEQ ID NO 916
                        Query Match
Best Local S
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Best Local
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LENGTH: 17
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APPLICANT: Gamper, Howard E
APPLICANT: Rice, Michael C.
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IIILE OF INVENTION: Stranded Oligonucleotides
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0.6%; Score 14.4;
il Similarity 93.8%; Pred. No. 46
15; Conservative
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93.8%; Pred. No. 4
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         Indels
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    Gaps
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; ORGANISM: Homo sapiens US-09-818-875-924
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                                                                                   SEQ ID NO 924
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-09-818-875-923
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Publication No. US20030051270A1
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PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
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PRIOR FILING DATE: 2000-10-30
NUMBER OF SEC ID NOS: 4385
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CURRENT APPLICATION NUMBER: US/09/818,875
CURRENT FILING DATE: 2001-03-27
CURRENT FILING DATE: 7001-03-27
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                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
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PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
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APPLICANT: Gamper, Howai
                                                                                                                               NUMBER OF SEQ ID NOS: 4385
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lication No.
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Rice, Michael C
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o. US20030051270A1
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Pred. No. 46;
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US-09-780-533A-2360
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APPLICANT: Chowrira, Bharat
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2360
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Best Local Similarity
Matches 15; Conserv
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                                                                        Query Match
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APPLICANT: Chowrira, Bharat
APPLICANT: Habbril, Pete
APPLICANT: Habbril, Pete
PILE OF INVENTION: Mcthod and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Blatt, Larry
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                                                                                                                              TYPE: RNA
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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                                                            Similarity
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Pred. No. 46
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US-10-060-756A-170
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Best Local (
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SEQ ID NO 170
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                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/327,898 PRIOR FILING DATE: 2001-10-09
                                                                                                                                                                                                                                                                                                                                    CURKENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
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LLE REFERENCE: PB0177
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mes 15; Conservative
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 09/864,761
FILING DATE: 2001-05-23
                                       APPLICATION NUMBER: US 60/327,898 FILING DATE: 2001-10-09
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 09/864,761
FILING DATE: 2001-05-23
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FILING DATE: 2001-01-30
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o. US20030046717A1
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Sequence Listing Engine
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Pred. No. 4
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; ORGANISM: Homo sapiens
US-10-060-756A-172
                                                                                                                   ; ORGANISM: Homo sapiens US-10-156-306-6015
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US-10-156-306-6015/c
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US-10-163-552-338
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LENGTH: 17
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Best Local 9
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CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
SOFTWARE: Patentin version 3.0
EQ ID NO 338
                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 8013
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
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APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of diseases
TITLE OF INVENTION: HER2
TITLE OF INVENTION: HER2
A (400/014)
                                                                                                                                               LENGTH: 17
TYPE: RNA
                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 17
TYPE: RNA
                                                                                                                                                                                                                                                          FILE REFERENCE: MBHB01-664-A
                                                                                                                                                                                                                                                                                                    PPLICANT: Ribozyme Pharmaceuticals, Inc
PPLICANT: McSwiggen, James
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                           1017 GGCCAGCACTCCCAGG 1032
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93.8%; Pred. No. 4
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; ORGANISM: Homo sapiens
US-10-209-787-915
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PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
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Publication No.
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PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
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CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
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CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
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ication No
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FILING DATE: 2000-03-27
APPLICATION NUMBER: US 60/208,538
FILING DATE: 2000-06-01
APPLICATION NUMBER: US 60/244,989
FILING DATE: 2000-10-30
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o. US20030217377A1
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93.8%; Pred. No. 4
    0.6%;
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         Score 14.4; DB 1;
Pred. No. 46;
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                                           Length 17;
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TITLE OF INVENTION: Stranded Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonuclectides
FILE REFERENCE: Napro-4
CURRENT ApplicATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
PRIOR APPLICATION NUMBER: US 9/818,875
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 6/192,176
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-10-30
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; ORGANISM: Homo sapiens
US-10-209-787-923
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PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 06/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
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SEQ ID NO 923
LENGTH: 17
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PRIOR FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 4385
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CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
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APPLICANT: Gamper, Howar
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D. US20030217377A1
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US-10-261-185-915
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; ORGANISM: Homo sapiens
US-10-209-787-924
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Best Local Similarity
Matches 15; Conserv
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LENGTH: 17
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Best Local Similarity
Matches 15; Conserv
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PRIOR APPLICATION NUMBER: PCT/US01/09761
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
                                                                                                                                                                 APPLICANT: Gamper, Howard B.
APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
FILE REFERENCE: NaPro-4CON
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SOFTWARE: Friedman macro Napro4
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PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR APPLICATION NUMBER: US 60/244,989
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PRIOR APPLICATION NUMBER: PCT/US01/09761
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-03-27
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CURRENT FILING DATE: 2002-09-27
PRIOR APPLICATION NUMBER: PCT/US01/09761
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APPLICANT: Rice, Michael C
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5. US20040014057A1
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Pred. No. 4
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Pred. No. 46;
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-261-185-923
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                                                                                                                           Sequence 924, Application US/10261185 Publication No. US20040014057A1
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SOFTWARE: Friedman macro Napro4
SEQ ID NO 916
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                                                    APPLICANT: Kmiec, Eric B.
APPLICANT: Gamper, Howar
APPLICANT: Rice, Michael
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonucleotides FILE REFERENCE: NaPro-4CON
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PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
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PRIOR APPLICATION NUMBER: PCT/US01/09761
PRIOR FILING DATE: 2001-03-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonucleotides FILE REFERENCE: NaPro-4CON
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PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
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TYPE: DNA
ORGANISM: Homo sapiens
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FILING DATE: 2000-03-27
APPLICATION NUMBER: US 60/192,179
FILING DATE: 2000-03-27
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Similarity 93.8%;
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                                                  Gamper, Howard B. Rice, Michael C.
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5. US20040014057A1
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Pred. No. 4
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Pred. No. 4
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CURRENT APPLICATION NUMBER: US/10/261,185

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; ORGANISM: Glycine max
US-09-969-373-2606
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; ORGANISM: Homo sapiens
US-10-261-185-924
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LENGTH: 17
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Best Local Similarity
                                                                                                                     GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
APPLICANT: Effertz, Roger J.
APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)A
CURRENT APPLICATION NUMBER: US/09/969,373
CURRENT FILING DATE: 2001-10-02
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PRIOR FILING DATE: 2001-05-15
NUMBER OF SEQ ID NOS: 4593
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CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US 09/754,853
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APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)
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                             PRIOR APPLICATION NUMBER: 60/251,015
PRIOR FILLING DATE: 2000-12-04
PRIOR APPLICATION NUMBER: 60/263,678
PRIOR FILLING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/273,037
PRIOR FILLING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 1579
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Best Local S
Matches 15
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PRIOR FILING DATE: 2001-01-13
PRIOR APPLICATION NUMBER: US 09/855,768
PRIOR FILING DATE: 2001-05-15
NUMBER OF SEQ ID NOS: 4593
SEQ ID NO 4139
LENGTH: 18
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SEQ ID NO 4141
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Patent No. US20020133852A1
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APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)A
CURRENT APPLICATION NUMBER: US/09/969,373
CURRENT FILING DATE: 2001-10-02
                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/005,956
CURRENT FILING DATE: 2001-12-03
                                                                                                                                                                                                APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: D0053NP
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PRIOR APPLICATION NUMBER: US 09/855,768
PRIOR FILING DATE: 2001-05-15
SOFTWARE: PatentIn version 3.0 EQ ID NO 715
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PRIOR FILING DATE: 2001-01-05
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ORGANISM: Glycine max
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Pred. No. 5
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Pred. No. 5
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CTGCCTTGGAGAAGAG

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CURRENT APPLICATION NUMBER: US/10/005,956

CURRENT FILING DATE: 2001-12-03

PRIOR APPLICATION NUMBER: 60/251,015

PRIOR FILING DATE: 2000-12-04

PRIOR APPLICATION NUMBER: 60/263,678

PRIOR APPLICATION NUMBER: 60/273,037

PRIOR APPLICATION NUMBER: 60/273,037

PRIOR APPLICATION NUMBER: 60/273,037

PRIOR FILING DATE: 2001-03-02

PRIOR FILING DATE: 2001-03-02

PRIOR FILING DATE: 2001-03-02

SOPTWARE: PATENTIN VETSION 3.0

SEQ ID NO 779

LENGTH: 18
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-005-956-715
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US-09-993-731-62
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; ORGANISM: Homo sapiens
US-10-005-956-779
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                 FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 62
SEQ ID NO 62
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                  Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                     Sequence 62, Applicate publication No. US200 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                      OTHER INFORMATION: Antisense Oligonucleotide -09-993-731-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: D0053NP
                                                                                                                                                                                                                                                                                                                                     APPLICANT: Brett P. Monia APPLICANT: Andrew T. Wat
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                                                                                                                                                          FEATURE:
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les 15; Conserv
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775 CTGCCTTTGAGAAGAG 790
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vo. US20030105040A1
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o. US20030113726A1
                                                0.6%; Score 14.4; illarity 93.8%; Pred. No. 73; Conservative 0; Mismatches
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Pred. No. 5
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Pred. No. 5
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RESULT 105
US-09-993-731-82
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Best Local S
Matches 16
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 82, Applicate Publication No. US20 GENERAL INFORMATION:
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Patent No. US20020048800A1
                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Wat
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APPLICANT:
APPLICANT:
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
                                                                                                                                                                                          APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
APPLICATION 10001-01-30
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
                                                                                                                                                        APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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                                               APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/US01/00667
                               APPLICATION NUMBER: PCT/US01/00661
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                                                                                                                                                                                                                                                                                                            LING DATE:
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INVENTION: MYOSIN-LIKE GENE EXPRESSED
SRENCE: AEOMICA-7
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No. US20030105040A1
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David R.
NUMBER: PCT/US01/00670
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Pred. No. 80;
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FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21

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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-2615
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SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 2616
LENGTH: 17
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CURRENT FILING DATE: 2001-05-25
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FITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                         PRIOR FILING DATE: 2001-02-05
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APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US01/00665
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APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/236,359
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                                                                                                                  FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
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APPLICATION NUMBER: PCT/US01/00661
                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US01/00662
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00664
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o. US20020048800A1
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PENN, Sharron G.
HANZEL, David K.
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Sequence 2368, Application US/09780533A

Publication No. US20030060611A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Habzyme Pharmaceuticals, Inc.
APPLICANT: Chowrira, Blatt, Larry
APPLICANT: MCSwiggen, Jim
APPLICANT: Habberii, Pete
FILE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00, 878-A (400/011)

CURRENT APPLICATION NUMBER: US/09/780,533A

CURRENT FILING DATE: 2001-02-09

PRIOR APPLICATION NUMBER: US 60/181,797

PRIOR FILING DATE: 2000-02-11

NUMBER OF SEQ ID NOS: 6679

SOFTWARE: Patentin version 3.0

SEQ ID NO 2368
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-780-533A-2096
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                                                                                 US-09-780-533A-2368
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CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Patentin version 3.0
SEQ ID NO 2096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 14; Conservative
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Best Local Similarity 100.0%; P
Matches 14; Conservative 0;
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Query Match 0.6%; Sometime 100.0%; Best Local Similarity 100.0%; Matches 14; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the
FILE REFERENCE: MBHSO, 878-A (400/01)
FILE REFERENCE: MBHSO, 878-A (400/01)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Blatt, Larry
                                                                                                    TYPE: RNA
ORGANISM: Homo sapiens
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on No. US20030060611A1
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100.0%; Pred. No.
ative 0; Mismatcl
                     0.6%; Score 14;
00.0%; Pred. No.
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Pred. No.
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                       DB 1;
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US-09-780-533A-2369
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Best Local :
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                                              PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR TILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00666
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PRIOR FILING DATE: 2000-05-26
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PLICANT: Haeberli, Pete
PLICANT: Haeberli, Pete
TLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
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Local L
14;
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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              CATION NUMBER: PCT/US01/00668
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20020048800A1
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EL, David K.
NUMBER: PCT/US01/00663
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RESULT 112
US-09-866-108-2593
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US-09-866-108-927
                       FILE REPERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILLING DATE: 2000-05-26
PRIOR FILLING DATE: 2000-10-04
PRIOR PELLONION NUMBER: US 60/236,359
PRIOR FILLING DATE: 2000-09-27
PRIOR PELLONION NUMBER: PCT/US01/00666
PRIOR FILLING DATE: 2001-01-30
PRIOR FILLING DATE: 2001-01-30
PRIOR FILLING DATE: 2001-01-30
PRIOR PELLONION NUMBER: PCT/US01/00667
PRIOR PELLONION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR PELLONION NUMBER: PCT/US01/00669
PRIOR PELLONION NUMBER: PCT/US01/00669
PRIOR PELLONION NUMBER: PCT/US01/00669
PRIOR PELLONION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00661
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Matches
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TYPE: DNA
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OR APPLICATION NUMBER: PCT/US01/00662
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00661
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00670
OR APPLICATION NUMBER: PCT/US01/00670
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: US 60/234,687
OR FILING DATE: 2000-09-21
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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RANK, David R.
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, Yonggang
INN, Sharron G.
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ON NUMBER: US 60/234,687
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Pred. No. 62;
0; Mismatches
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片
; ORGANISM: Homo US-09-866-108-6611
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Best Local Similarity
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                                                                      SEQ ID NO 6611
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ORGANISM: Homo sapiens
-09-866-108-2593
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TITLE OF I
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NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
EQ ID NO 2593
LENGTH: 17
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PRIOR FILING DATE: 2000-05 07
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CURRENT FILING DATE: 2001-05-25
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                                                                                                       PRIOR FILING DATE: 2001-02-
NUMBER OF SEQ ID NOS: 15752
                                                                                                                             PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
                                   TYPE: DNA
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                                                    ENGTH:
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FILING DATE: 2000-09-27
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APPLICATION NUMBER: PCT/US01/00661
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                           CATION NUMBER: PCT/US01/00669
                                                                                                                                                                                                                                                                                                                                                                                                          CATION NUMBER: PCT/US01/00665
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NN, Sharron G.
NZEL, David K.
                   sapiens
                                                                                          Sequence Listing Engine
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Pred. No. 62;
0; Mismatches
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Query Match

0.5%;

Score 13.8;

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Length 17;

RESULT 115 US-09-866-108-7854/c

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; ORGANISM: Homo sapiens US-09-866-108-6612
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                                                                             Query Match
Best Local
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SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 66612
LENGTH: 17
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PRIOR FILING DATE: 2000-10-04
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PRIOR FILING DATE: 2000-05-26
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APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MATK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART
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PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
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                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00669
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                              1221 CAGAACCTCCAGCATGT 1237
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o. US20020048800A1
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                                                                                 Score 13.8;
Pred. No. 6;
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RESULT 116
US-09-866-108-7855/c
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US-09-866-108-7854
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SOFTWARE: Aecmica Sequence Listing Engine
SEQ ID NO 7854
LENGTH: 17
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Best Local Similarity 88.2%;
Matches 15; Conservative
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                                                                                                                                                            Sequence 7855, Application US/09866108 Patent No. US20020048800A1
                                                                              APPLICANT:
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
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FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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PRIOR APPLICATION NUMBER: US 60/236,359
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PRIOR FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/266,860
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JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
SHANNON, Mark
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N, Sharron
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Pred. No. 62;
0; Mismatches
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-7855
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                Patent No. US2003
CURRENT APPLICATION NUMBER: US/09/866,108 CURRENT FILING DATE: 2001-05-25 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26 PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR PILING DATE: 2000-09-27
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PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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CURRENT FILING DATE: 2001-05-25
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APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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Pred. No. 6
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; ORGANISM: Homo sapiens
US-09-866-108-8082
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SEQ ID NO 8082
LENGTH: 17
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CURRENT FILING DATE: 2001-05-25
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                                                                                                                                                                                                                                                                                                                                FILE REFERENCE:
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                                                                                                                                                                                                                                                                                                                                             PLICANT: SHANNON, Mark
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
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                                                                                                                            FILING DATE: 2000-09-2/US01/00666
                                                                                                                                                                             APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
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APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
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APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-
                                                                                   FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
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APPLICATION NUMBER: PCT/US01/00661
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                                                  APPLICATION NUMBER: PCT/US01/00669
                                                                      FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
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c. US20020048800A1
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HANZEL, David K
RANK, David R.
CHEN, Wensheng
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: 2001-01-30
NUMBER: PCT/US01/00665
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88.2%;
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Pred. No. 62;
0; Mismatches
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; ORGANISM: Homo sapiens
US-09-866-108-8648
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PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engir
SEQ ID NO 8648
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 10738, Application US/09866108 Patent No. US20020048800A1 GENERAL INFORMATION:
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Best Local Similarity 88.2%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggan
APPLICANT: PENN, Shar
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OR APPLICATION NUMBER: PCT/US01/00663
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00662
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00661
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00670
OR APPLICATION NUMBER: PCT/US01/00670
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00662
OR PILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00661
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00670
OR APPLICATION NUMBER: PCT/US01/00670
OR FILING DATE: 2001-01-30
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                                                                                                                                                         APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
                                                                                                                                                                                                                          APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00665 FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00667
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Pred. No. 62;
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APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21

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Best Local :
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LENGTH: 17
                                                        SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 10740
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PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
                                                                                                                       PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION UNMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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NUMBER OF SEQ ID NOS: 15752
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ORGANISM: Homo sapiens
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les 15; Conserv
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APPLICATION NUMBER: PCT/US01/00664
PTITING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
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LING DATE: 2001-01-30
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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ilarity 88.2%;
Conservative
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Pred. No. 62;
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US-09-864-785-1560/c

Sequence 1560, Application US/09864785 Patent No. JUS20020177568A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Stinchcomb, Dan APPLICANT: Draper, Ken

RESULT 122

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SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-860-784-28
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Best Local
                                                                           Query Match
Best Local
                                                            Matches
                                                                                                                                                                                                               TELEX: 904136
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/860,784
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FILING DATE: 04-APR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UHLMANN, EUGEN TITLE OF INVENTION: G CAP-STABILIZED OLIGONUCLEOTIDES NUMBER OF SEQUENCES: 105
                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                            1775 GGAGGAGGCGGAGGAGG 1791
                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
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15; Conserv
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                                                                                                                                                                  LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
GGAGGATGCTGAGGAGG 17
                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 31,298
REFERENCE/DOCKET NUMBER: 18748/264/HOCE
                                                                                                                                                                                                                                                                                                                             NAME: SANDERCOCK, COL
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 21-May-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Washington
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Pred. No. 6
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; ORGANISM: Homo sapiens
US-09-825-805-667
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                                                                                                                                                                                              SOFTWARE: Pate
SEQ ID NO 667
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sweedler, Dave
APPLICANT: Zinnen, Shawn
TITLE OFT Zinnen, Shawn
TITLE OFT SINCEN, Nucleotide Triphosphate and their Incorporation into Oligonucleol
FILE REFERENCE: MEHBOO-831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILING DATE: 2001-09-27
CURRENT FILING DATE: 2001-09-27
                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 1558
                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/578,223
PRIOR TILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 09/476,387
PRIOR FILING DATE: 1999-12-30
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TYPE: RNA
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                                                                                        Local
                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 09/474,432
FILING DATE: 1999-12-29
APPLICATION NUMBER: 09/301,511
FILING DATE: 1999-04-28
APPLICATION NUMBER: 09/186,675
FILING DATE: 1998-11-04
                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/083,727 FILING DATE: 1998-04-29
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                                    1268 GGAAGAGGCTGAGGGCA 1284
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                                                                                        Similarity
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Beaudry, Amber
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                                                                        Conservative
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Pred. No. 62;
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CURRENT FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR TILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
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SEQ ID NO 927
LENGTH: 17
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SOFTWARE: Friedman macro N
SEQ ID NO 928
                                                                                  Matches
                                                                                                     Query Match
Best Local
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APPLICANT: Kmiec, Eric B.
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Best Local
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PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
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APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
                                                                                                                                                                                                     LENGTH: 17
TYPE: DNA
                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/244,989 PRIOR FILING DATE: 2000-10-30
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PRIOR APPLICATION NUMBER: US 60/192,176
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APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
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                                                                                                                                                                                 ORGANISM: Homo sapiens
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URRENT APPLICATION NUMBER: US/09/818,875
URRENT FILING DATE: 2001-03-27
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                                       1264 AGCTGGAAGAGGCTGAG 1280
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  AGCTGGAAGAGTCTGGG
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o. US20030051270A1
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; Pred. No. 62;
0; Mismatches
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Pred. No. 6
                                                                                    Mismatches
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RESULT 128
US-09-780-533A-2361
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US-09-780-533A-2358
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; ORGANISM: Homo sapiens
US-09-780-533A-899
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Sequence 2361, Application US/09780533A Publication No. US20030060611A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 15; Conserv
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Best Local Similarity 88.2%;
Matches 15; Conservative
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Publication No.
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APPLICANT: Chownira, Bharat
APPLICANT: Chownira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MEHBOO,878-A (400/011)
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR TILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Patentin version 3.0
SEQ ID NO 2358
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TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
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APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
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Chowrira, Bharat
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88.2%;
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Pred. No. 6;
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Pred. No. 6
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-780-533A-2361
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US-09-780-533A-2363
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PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOPTWARE: Patentin version 3.0
SEQ ID NO 2361
                                                                                                                       Sequence 2363, Application US/09780533A
Publication No. US20030060611A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 2362
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Best Local Similarity 88.2
Matches 15; Conservative
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Best Local Similarity 88.2%;
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APPLICANT: Chowrira, Bharat
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberii, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Blatt, Larry
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APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
                        FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENGTH:
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P: McSwiggen, Jim
I: Chowrira, Bharat
I: Haeberli, Pete
INVENTION. Method and Reagent for the Inhibition of NOGO Gene
ERENCE: MBHB00,878-A (400/011)
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o. US20030060611A1
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Pred. No. 6
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US-09-780-533A-2462
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US-09-848-754A-2225
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US-09-848-754A-2225
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SOFTWARE: Patentin version 3.0
SEQ ID NO 2363
LENGTO
                                                            APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MSHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2225
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Matches
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 2462
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Best Local Similarity
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APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
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CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
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ORGANISM: Homo sapiens
                ORGANISM: Homo sapiens
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Pred. No. 62
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Query Match
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US-09-930-423-189
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                                                                                                                          ; ORGANISM: Homo Sapiens
US-09-930-423-795
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 189
LENGTH: 17
                                                                                                                                                                                    APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MEHB00, 918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTMARE: PatentIn version 3.0
SEQ ID NO 795
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APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
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Best Local Similarity
Matches 15; Conserv
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TITLE OF INVENTION: Method and Reagent for
FILE REFERENCE: MeBBO0,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
                                                                                                                                                         LENGTH: 17
TYPE: RNA
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GCGCAGATGGCAGCCAT 1
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Pred. No. 62;
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Pred. No. 62;
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RESULT 135 US-09-930-423-1533/c

Sequence 1533, Application US/09930423 Publication No. US20030092003A1 GENERAL INFORMATION: APPLICANT: Ribozyme Pharmaceuticals, Inc.

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US-09-930-423-1533
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CURRENT APPLICATION NUMBER: US/09/930,423

CURRENT FILING DATE: 2001-08-15

NUMBER OF SEQ ID NOS: 4553

SOFTWARE: Patentin version 3.0

SEQ ID NO 1533

LENGTH: 17
                                                      FILE REFERENCE: RFI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
SEQ ID NO 1895
LENGTH: 17
                                                                                                                                                                                    Sequence 1895, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 244, Application US/09740332
publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
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Best Local Similarity
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CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
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NAME/KEY: misc_feature
LOCATION:
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FEATURE:
                  ORGANISM: artificial sequence
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Similarity 88.2%;
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Pred. No. 6
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Pred. No. 6;
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; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-4311
                                                                                                                                               NUMBER OF SEQ ID NOS: 9704
SEQ ID NO 4311
LENGTH: 17
TYPE: RNA
ORGANISM: artificial sequence
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US-09-740-332-2660/c
                                                                                                                                                                                                                                                                                 Sequence 4311, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Related
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
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Publication No. US20
GENERAL INFORMATION:
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Best Local Similarity 64.7
Matches 11; Conservative
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TYPE: RNA
ORGANISM: artificial sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION:
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CURRENT FILING DATE: 2001-03-26
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TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
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SOFTWARE: PatentIn version 3.0
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NAME/KEY: misc_feature
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Local Similarity
les 15; Conserv
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88.2%; Pred. No. 62;
tive 0; Mismatches
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              Score 13.8;
Pred. No. 62;
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Pred. No. 62;
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US-09-745-237A-795/c
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US-09-745-237A-189/c
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LENGTH: 17
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                                                                                                                            Sequence 1533, Application US/09745237A Publication No. US20030143708A1
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Best Local
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APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING_DATE: 2002-04-15
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TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (NEHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 455
NUMBER OF SEQ ID NOS: 455
NUMBER OF SEQ ID NOS: 455
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                                                                                      APPLICANT: Ribozyme Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Homo sapiens
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Local Similarity 88.2%;
es 15; Conservation
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: McSwiggen, Jim
INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
!RENCE: 400/007 (MBHB00-918-A)
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o. US20030143708A1
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5. US20030143708A1
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88.2%;
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Pred. No. 62;
0; Mismatches
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Pred. No. 6
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; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-1895
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; ORGANISM: Homo sapiens
US-09-745-237A-1533
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Best Local S
Matches 15
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TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
SEQ ID NO 1895
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APPLICANT: R.bozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment
TITLE OF INVENTION: Hepartitis C Virus Infection
FILE REFERENCE: MBH300-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
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CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1533
LENGTH: 17
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Best Local Similarity
Matches 15; Conserv
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Publication No.
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LOCATION:
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                                                   NAME/KEY: misc_feature
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; OTHER INFORMATION: oligonucleotide substrate
US-09-817-879-2660
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US-09-817-879-4311
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Best Local Similarity
Matches 15; Conserv
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CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
SEQ ID NO 2660
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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
                                                                                                                                                                                                                                   SEQ ID NO 4311
LENGTH: 17
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CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTMARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                           APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepartitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
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hblication No. US20030171311A1
HENERAL INFORMATION:
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TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
                                                                                                                              LOCATION:
OTHER INFORMATION: oligonucleotide substrate
                                                                                                                                                                                      ORGANISM: artificial sequence FEATURE:
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NAME/KEY: misc_feature
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TYPE: RNA
ORGANISM: artificial sequence
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Pred. No. 6
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Pred. No. 62;
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Pred. No. 62;
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APPLICANT: MCSWiggen, Jim
APPLICANT: MCSWiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of
TITLE OF INVENTION: HER2
FILE REFERENCE: MBHBOI-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
SOFTMARE: Patentin version 3.0
SEQ ID NO 337
LENGTH: 17
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-060-756A-358
                                                                                                                                        US-10-163-552-337
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Publication No. US2
GENERAL INFORMATION
                                                                                     Query Match
Best Local
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                                                                    Matches
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Publication No. US20030105051A1
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CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
                                                                                                                                                                      LENGTH: 17
TYPE: RNA
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PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                       ORGANISM: Homo sapiens
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                                                                                 Local
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 09/864,761
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                                1268 GGAAGAGGCTGAGGGCA 1284
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l Similarity 88.2%;
15; Conservation
                                                                    l Similarity 88.2
15; Conservative
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o. US20030046717A1
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                                                                                   88.2%;
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Pred. No. 62;
0; Mismatches
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                                                                                     Score 13.8;
Pred. No. 6
                                                                      Mismatches
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                                                                                                    Length: 17;
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RESULT 150
US-10-156-306-5922
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Best Local S
Matches 13
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 5905
LENGTH: 17
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GENERAL INFORMATION:
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases (
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR APPLICATION NUMBER: PCT/US 02/16840
PRIOR PRIOR PRIOR PRIOR FILING DATE: 2002-05-29
PRIOR PRIOR FILING DATE: 2001-09-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McSwiggen, James
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTWARE: Patentin version 3.0
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APPLICANT: McSwiggen, James
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
TILE OF INVENTION: Levels of IKK-Gamma and PKR
TILE REPERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
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APPLICANT: McSwiggen, James
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ORGANISM: Homo sapiens
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Local Similarity 88.2%;
les 15; Conservation
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13; Conserv
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Pred. No. 6
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                            PRIOR APPLICATION NUMBER: US 09/818,875
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/2044,989
PRIOR APPLICATION NUMBER: US 60/244,989
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PRIOR FILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 721
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 927, Applic Publication No. US20 GENERAL INFORMATION:
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Publication No.
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 3391
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Best Local Similarity 88.2%;
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APPLICANT: Fosnaugh, Kathy
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR
FILE REFERENCE: 400/056 (MBHB01-1110)
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
                                                                                                                                                                                                                     APPLICANT: Kmiec, Eric B.
APPLICANT: Gamper, Howard B.
APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonucleotides
FILE REFERENCE: Napro-4
FULS REFERENCE: Napro-4
CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
CURRENT FILING DATE: 2002-07-30
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
               RIOR FILING DATE: 2000-10-30
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o. US20030191077A1
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ID NOS: 4385
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Pred. No. 6;
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Pred. No. 6:
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-209-787-927
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US-10-209-787-928
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Best Local Similarity
Matches 15; Conserv
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LENGTH: 17
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Best Local
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                                                                                                                                     TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonucleotides
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PRIOR APPLICATION NUMBER: US 09/818,875
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-03-27
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                                                        CURRENT APPLICATION NUMBER: US/10/261,185
CURRENT FILING DATE: 2002-09-27
CRICK APPLICATION NUMBER: PCT/US01/09761
PRIOR FILING DATE: 2001-03-27
                                                                                                                                                                                                   ENERAL INFORMATION:
APPLICANT: Knice, Eric B.
APPLICANT: Gamper, Howard E
APPLICANT: Rice, Michael C.
                                                                                                                                                                                                                                                                                 equence 927, Aublication No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PPLICANT: Kmiec, Eric B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        blication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRICANT: Gamper, Howard B.

PRICANT: Rice, Michael C.

TLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single

TLE OF INVENTION: Stranded Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/192,179
FILING DATE: 2000-03-27
APPLICATION NUMBER: US 60/208,538
FILING DATE: 2000-06-01
APPLICATION NUMBER: US 60/244,989
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                   APPLICATION NUMBER: US 60/192,176 FILING DATE: 2000-03-27
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APPLICATION NUMBER: US 60/192,179
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Similarity 88.2%;
                                                                                                                                                                                                                                                                                    Application US/10261185
o. US20040014057A1
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                                                                                                                                                                                                                      Howard B.
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Pred. No. 6
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US-10-261-185-927
                                                                                                                                                                                                                                                                                                                                                                                          US-10-261-185-928
                                                                                                          Sequence 72, Applica Publication No. US20 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Best Local
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CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
                                APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 17
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 4385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 2002-09-27
PRIOR APPLICATION NUMBER: PCT/US01/09761
PRIOR FILING DATE: 2001-03-27
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APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
TILE REFERENCE: NaPro-4CON
CURRENT APPLICATION NUMBER: US/10/261,185
CURRENT FILING DATE: 2002-09-27
CURRENT FILING DATE: 2002-09-27
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NUMBER OF SEQ ID NOS: 4385
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                                                                        APPLICANT: Brett
APPLICANT: Andre
                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                      Local
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FILING DATE: 2000-03-27
APPLICATION NUMBER: US 60/192,179
FILING DATE: 2000-03-27
APPLICATION NUMBER: US 60/208,538
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FILING DATE: 2000-06-01
APPLICATION NUMBER: US 60/244,989
FILING DATE: 2000-10-30
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1 Similarity 88.2%;
15; Conservation
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                                                                                                                                              Application US/09993731
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o. US20040014057A1
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Pred. No. 6
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                                                        INHIBITOR-KAPPA B-R EXPRESSION
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RESULT 159
US-10-056-414-33/c
/Sequence 33, Application US/10056414
/ Publication No. US20030003469A1
/ Publication INFORMATION:
GENERAL INFORMATION:
GENERAL STAND: Stinchcomb, Dan T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Oligonucleotide US-09-880-313A-235
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CURRENT FILING DATE: 2001-06-13
NUMBER OF SEQ ID NOS: 276
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 235
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Best Local Similarity
Matches 14; Conserv
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APPLICANT: Flemington, Erik K
TITLE OF INVENTION: Adaptors and Methods of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 235, Application US/09880313A
Publication No. US20030044791A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
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                                            ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1247 GGTCCGGCTGCAGCA 1261
                                                                                                                                                                                                                                              McSwiggen, James
TITLE OF INVENTION: RIBOZYME TREATMENT OF
DISEASES OR CONDITIONS
RELATED TO LEVELS OF
                                                                                                                                                                                                               NUMBER OF SEQUENCES: 830
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15; Conserv
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COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C.
                                                                                                               CITY: Los Angeles
STATE: California
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Pred. No. 96;
0; Mismatches
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Pred. No. 52;
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                                                 1.44 Mb
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                                                                           RESULT 161
US-10-314-405-43
                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens
US-10-163-552-1980
                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
SOFTWARE: Patentin version 3.0
SEQ ID NO 1980
LENGTH: 15
TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1980, Application US/10163552
Publication No. US20030105051A1
Sequence 43, Application US/10314405
publication No. US20030108940A1
GENERAL INFORMATION:
APPLICANT: Hidetoshi, Inoko
                                                                                                                                                                                                                      Query Match 0.1
Best Local Similarity 93.
Matches 14; Conservative
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Nucleic acid treatment TITLE OF INVENTION: HER2
FILE REFERENCE: MEHBO1-1653-A (400/014)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
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INFORMATION FOR SEQ ID NO: 33:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RETURNAL: Marburg, Richard ...
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/157
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/056,414
FILING DATE: 23-Jan-2002
CLASSIFICATION: Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 33:
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                                                                                                                                                                                       1267 TGGAAGAGGCTGAGG 1281
                                                                                                                                                   15 TGGAAGACGCTGAGG 1
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/291,932A FILING DATE: August 15, 1994 APPLICATION NUMBER: 08/245,466 FILING DATE: May 18, 1994 APPLICATION NUMBER: 07/987,132 FILING DATE: December 7, 1992
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Pred. No. 52;
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Pred. No. 52;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of diseases or conditions related to level
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RESULT 163
US-10-156-306-7883/c
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; ORGANISM: Homo sapiens
US-10-156-306-7883
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US-10-314-405-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 7867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 7867, Application US/10156306 Publication No. US20030119017A1 GENERAL INFORMATION:
                                                                              SOFTWARE: PatentIn version 3.0 SEQ ID NO 7883
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LENGTH: 15
                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                      CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
                                                                                                                                                                             APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, James TITLE OF INVENTION: Enzymatic Nucleic Acid TITLE OF INVENTION: Levels of IKK-Gamma an FILE REFERENCE: MBHB01-664-A (400/050)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Levels of IKK-Gamma and PKR FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
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APPLICANT: McSwiggen, James

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate

TITLE OF INVENTION: Levels of IKK-Gamma and PKR
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CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: US/09/713,616
PRIOR FILING DATE: 2000-11-15
NUMBER OF SEQ ID NOS: 46
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                          equence 7883, Application US/10156306
ublication No. US20030119017A1
                                       TYPE: RNA
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                                                           ENGTH:
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ITLE OF INVENTION: NOVEL PO
ILE REFERENCE: 06501-069001
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les 13; Conserv
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86.7%; Pred. No. 5
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Pred. No. 5
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                                                                                                                                                                                                                         Treatment of Diseases or Conditions Relat
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US-10-043-875-406/c
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CURRENT APPLICATION NUMBER: US/10/428,826
CURRENT FILING DATE: 2003-05-05
PRIOR APPLICATION NUMBER: US/09/601,326
PRIOR APPLICATION NUMBER: PCT/US99/02630
PRIOR APPLICATION NUMBER: US 09/019,793
PRIOR FILING DATE: 1999-04-19
PRIOR APPLICATION NUMBER: US 09/019,793
PRIOR FILING DATE: 1998-02-06
PRIOR FILING DATE: 1998-02-06
                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: De Smet, Koenraad
APPLICANT: Stuyver, Lieven
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APPLICAM: Stuyer, Lieven
APPLICAM: Stuyer, Lieven
TITLE OF INVENTION: Method for Detection of Drug-Induced Mutations i
TITLE OF INVENTION: Transcriptase Gene
FILE REFERENCE: 11362-0033-NPUG01 (INNS:033)
CURRENT APPLICATION NUMBER: US/10/043,875
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 60/286,102
PRIOR FILING DATE: 2001-04-24
PRIOR FILING DATE: 2001-04-24
PRIOR FILING DATE: 2001-04-26
PRIOR FILING DATE: 2001-04-27
PRIOR APPLICATION NUMBER: EP 01870085.6
PRIOR FILING DATE: 2001-04-20
PRIOR APPLICATION NUMBER: EP 01870095.4
PRIOR FILING DATE: 2001-01-11
                                                                                                                                                                                                                                                                                                                                            Sequence 406, Application US/10043875
Publication No. US20030054339A1
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Best Local
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Best Local
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GENERAL INFORMATION:
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APPLICANT: ZHANG, YANJIN

TITLE OF INVENTION: PROTEINS ENCODED BY POLYNUCLEIC ACIDS OF PORCINE

TITLE OF INVENTION: REPRODUCTIVE AND REPIRATORY SYNDROME VIRUS (PPRSV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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FILING DATE: 1994-09-01
APPLICATION NUMBER: US 08/131,625
FILING DATE: 1993-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/478,316 FILING DATE: 1995-06-07
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14; Conser
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. US20030186225A1
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Pred. No. 52;
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; OTHER INFORMATION: V92M Val probe
US-10-108-732-58
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US-10-108-732-58
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; ORGANISM: Human immunodeficiency virus
US-10-043-875-406
                                                                                                                                                                                                                                                                                       RESULT 167
                                                                                                                                                                                                                                                                       US-10-043-875-407/c
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Publication No
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Best Local Similarity 100.0%;
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Publication No. US20030054339A1
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             APPLICANT: De Smet, Koenraad
APPLICANT: Stuyver, Lieven
TITLE OF INVENTION: Method for Detection of Drug-Induced Mutations in the HIV Reverse
TITLE OF INVENTION: Transcriptase Gene
FILE REFERENCE: 11362-0033-NFUSO1 (INNS:333)
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 05/286,102
PRIOR APPLICATION NUMBER: 60/286,102
PRIOR APPLICATION NUMBER: BP 01070085.6
PRIOR FILING DATE: 2001-04-20
PRIOR FILING DATE: 2001-04-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/108,732
CURRENT FILING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: US 60/279,515
PRIOR FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 76
SOFTWARE: Patentin version 3.1
EQ ID NO 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Sturm, Richard A
APPLICANT: Gruis, Nelleke A
APPLICANT: Van Der Velden, Pieter
APPLICANT: Bergman, Wilma
APPLICANT: Frants, Rune R
TITLE OF INVENTION: MELANOMA RISK DETECTION
FILE REFERENCE: 8795-27U1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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APPLICATION NUMBER: EP 01870005.4
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Martin, Nicholas G
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10. US20030175721A1
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Conservative 0;
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; Pred. No.
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TYPE: DIA
; ORGANISM: Human immunodeficiency virus
US-10-043-875-407
                                             FILE REFERENCE: 2702.1001-011
CURRENT APPLICATION NUMBER: US/10/150,510
CURRENT FILING DATE: 2002-08-23
PRIOR APPLICATION NUMBER: 09/999,031
PRIOR FILING DATE: 2001-11-01
PRIOR APPLICATION NUMBER: PCT/US00/12179
PRIOR FILING DATE: 2000-05-04
PRIOR APPLICATION NUMBER: 60/133,224
PRIOR FILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 10
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PRIOR FILING DATE: 2000-05-04
PRIOR PELICATION NUMBER: 60/133,224
PRIOR FILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 3
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APPLICANT: Garchon, Henri-Jean
TITLE OF INVENTION: DIAGNOSIS OF GLAUCOMA
FILE REFERENCE: 2702.1001-004
FULE REFERENCE: 2702.1001-004
CURRENT APPLICATION NUMBER: US/09/999,031A
CURRENT FILING DATE: 2002-04-11
CURRENT FILING DATE: 2002-04-11
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SOFTWARE: Patentin version 3.1
SEQ ID NO 407
LENGTH: 16
              SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 3
                                                                                                                                                                                                                                                                                      Sequence 3, Application US/10150510 Publication No. US20030068632A1 GENERAL INFORMATION:
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Patent No. US20020164608A1
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Best Local .
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Best Local Similarity
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                                                                                                                                                                                                                              APPLICANT: Garchon, Henri-Jean
TITLE OF INVENTION: DIAGNOSIS OF GLAUCOMA
FILE REFERENCE: 2702.1001-011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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TH: 16
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87.5%;
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00.0%; Pred. No.
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Pred. No. 8
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; LOCATION: (1)...(16)
; OTHER INFORMATION: Oligonucleotide
US-10-150-510-3
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US-09-866-108-8648/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: ANTISENSE COMPOSITIONS TARGETED TO BETA1-ADRENOCEPTOR-SPECIFIC TITLE OF INVENTION: METHODS OF USE FILE REFERENCE: 4300.013900 CURRENT APPLICATION NUMBER: US/10/308,503 CURRENT FILING DATE: 2003-12-03
              CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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PRIOR FILING DATE: 1998-09-14
PRIOR APPLICATION NUMBER: PCT/US99/21007
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APPLICANT: JI, Yongga
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APPLICANT: ZHANG, YI
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ORGANISM: Artificial Sequence
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LENGTH: 16
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Local Similarity 87.5%;
                                                                                  REFERENCE: AEOMICA-
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                                                                                                                                                                                                                                                                                                                                                                                           1649 TGCCCAGCTGCAGAGG 1664
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                                                                                                                                                                                                                                     INFORMATION
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                                                                                         T: CHEN, Wensheng
T: SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                                                                                                 RANK, David R.
CHEN, Wenshor
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b. US20030191080A1
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GB 24263.6
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Pred. No. 8
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Pred. No. 8
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                                                                                                                      ) OTHER INFORMATION: Antisense Oligonucleotide US-09-993-731-61
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US-09-866-108-8648
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SEQ ID NO 8648
LENGTH: 17
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                                                                        Query Match
Best Local
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LENGTH: 20
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Best Local Similarity
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PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomics Of
                                                                                                                                                                                                                             FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                          APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                      FEATURE:
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APPLICATION NUMBER: US 60/234,687
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APPLICATION NUMBER: PCT/US01/00666
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                            777 GCCTTTGAGAAGAGCT 792
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 GCCTTGGAGAAGAGGT 16
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                                                             Score 12.8; I
Pred. No. 1.5e
0; Mismatches
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Pred. No. 9
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US-09-93-731-77

Sequence 77, Application US/09993731

Publication No. US20030105040A1

GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
APPLICANT: ANTIENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
TYPE: DNA
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TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
CURRENT APPLICATION NUMBER: US/09/993,731
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 63
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
COTHER IMPORMATION: Antisense Oligonucleotide
US-09-993-731-63
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US-09-993-731-74
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US-09-993-731-63
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CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 74
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Brett P. Monia
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF INHIBITOR-KAPPA B-R EXPRESSION
FILE REFERENCE: RTS-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
nes 15; Conserv
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Search completed: April Job time : 5 secs
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                                                                                                                                                                        ; ORGANISM: Artificial Sequence , FEATURE: OTHER INFORMATION: Antisense Oligonucleotide US-09-993-731-77
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Best Local Similarity 78.9
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                                                                                  1866 GGCCTGACCCCGCAGCTGG 1884
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1M0079M16R Mouse 10kb plasmid
                            clone UUGC1M0079M16 R, AZ345323
AZ345323 GI:10424560
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musculus
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Copyright (c) 1993 - 2004 Compus
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                                                           27 bp DNA linear GSS 29-SEP-200
smid UUGCIM library Mus musculus genomic
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ACCESSION: AZ321746
ACCESSION: AZ775540
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ACCESSION: AZ321746
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RESULT 2
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                                                          AZ321746 21 bp DNA linear GSS 29-SEP-200 IM0042N20F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0042N20 F, genomic survey sequence.

AZ321746 AZ321746.1 GI:10374795 GSS.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
1 (bases to 27)
Mus musculus (house mouse)
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil Islam, H., Longacre, S., Mahmoud, N., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         308,
                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: pWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                 Score 21.8; DB Pred. No. 0.64; 0; Mismatches
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        Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 95.0
                                                                                                                                                                                                                                                                                                                                                                                                     1772
                                                                                                                                             23 bp DNA linear GSS 13-DEC-20(1M0435A21F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0435A21 F, genomic survey sequence.
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Mouse whole genome scaffolding with paired end reads from 10kb
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                High quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
Contact: Robert B.
                                          Mus musculus
                                                                        Mus
                                                                                                     SSD
                                                                                                                          AZ610186.1 GI:11732376
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Insert Length: 10000 Std Error:
Plate: 0042 row: N column: 20
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
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                                                                        musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Nouse DNA Resources (documents/dnares). The DNA (http://www.jax.org/resources/documents/dnares). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNP42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse. DNA was annealed to adaptored vector DNA, and transformed into adaptored for ampicillin resistance."
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PW021Nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
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/strain="C57BL/6J"
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Class: plasmid ends
Class: plasmid ends
High quality sequence stop: 23.
Location/Qualifiers
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Nouse whole genome scaffolding with paired end reads from 10kb
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 19)
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                    Mus musculus (house
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AZ775540.1
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Insert Length: 10000 Std Error:
Plate: 0435 row: A column: 21
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                                                                             AI564678 Inear EST 14-MAY-1 tq78g03.x1 NCI CGAP_Utl Homo sapiens cDNA clone IMAGE:2214964 3' similar to TR:Q15214 Q15214 SAILVARY PROLINE-RICH PROTEIN 1 contains element MSR1 repetitive element ;, mENA sequence.
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Lengrh: 10000 Std Error:
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 16)
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
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/clone="UUGC2M0008H15"
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strain="C57BL/6J"
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Mouse whole genome scaffolding with paired end reads from 10kb
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
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DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                 University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                    Contact: Robert
                                                                                                                                                                                        Unpublished
                                                                                                                                                                                                          plasmid inserts
                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus (house mouse)
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National Cancer Institute, Cancer Genome Anat
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USA
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/lab host="DHIOD"
/lab host="DHIOD"
/clone_lib="NCI_CGAP_Ut1"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall;
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Sycrope_insert size 1.75 kb. Life Technologies catalog #:
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/mol type="mtNA"
/db_xref="taxon:9606"
/clone="IMAGB:2214964"
/tissue type="well-differentiated endometrial
/denocarcinoma, 7 pooled tumors"
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0; Mismatches
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RESULT 7
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17; Conserv
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Seq primer: CACACAGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 27.
Location/Qualifiers
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
                                                                                                                 Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@gbbo.com, bhnahm@bio.myongji.ac.kr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HDA1--04-H13.gl OsHDAC1-overexpressing transgenic rice lambda ph cDNA library I (HDA1) Oryza sativa cDNA clone HDA1--04-H13, mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Large-scale Sequencing Analysis Unpublished (2003)
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/organism="Oryza sativa"
/mol_type="mRNA"
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/strain="C57BL/6J"
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'clone="UUGC1M0079M16"
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Pred. No. 11;
0; Mismatches 9;
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RESULT 9
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CF332055 12 bp mRNA linear EST NACL--08-G14.g1 Rice callus plasmid cDNA library (NACL) sativa cDNA clone NACL--08-G14, mRNA sequence.
                                                                                                                                                                                                        12;
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Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14 bp mRNA linear EST 17 5009-0-68-B01.t.1 Chilcoat/Turkewitz cDNA (large fraction) Terrabymena thermophila cDNA, mRNA sequence.
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Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E.,
Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila, strain CU428.1
Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Chicago
920 E. 58th Street, Chicago,
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Molecular Genetics and Cell Biology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /dev stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="p.coli_SOLR"
/clone_lib="0sHDAC1-overexpressing transgenic rice lambda
phage_CDNA_library_I_(RDA1)"
/note="Vector: pBluescript_SK(+); Site_1: EcoRI; Site_2:
/note="Vector: pBluescript_SK(+); Site_1: EcoRI; Site_2:
XhoI; Callus was treated with ABA(20um) for lhour. cDNA
was inserted into lambda Uni-ZAP_XR vector at 5' end with
BCORI and 3' end with XhoI site. mRNA was derived from
                                                                                                                                                                                                                                                                          /db_xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (20
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                                                                                                                                                                                                                                                                                          /organism="Tetrahymena thermophila"
/mol_type="mRNA"
/strain="CU428.1"
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/db_xref="taxon:4530"
/clone="HDA1--04-H13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ocation/Qualifiers
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Pred. No. 5;
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                                         EST 18-AUG-2003
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Matches 11; Conserv
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BQ593629 13 bp mRNA linear EST 06-DEC-2002
E012766-024-026-N02-SP6_MPIZ-ADIS-024-developing root Beta vulgaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
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Oryza sativa
                                                  Contact: Weisshaar B
ADIS DNA core facility at MPIZ
ANAX-Planck-Innstitute for Plant Breeding Research
Carl-von-Linne 10, 50829 Koeln, Germany
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Genomics and Genetics Institute, Greendene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
Fax: 00492215062851
Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 13 Std Error: 0.00
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Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet general J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                 Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.

1 (bases 1 to 13)
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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CF332055.1
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/clone="NACL--08-G14"
/tissue type="callus"
/dev_stage="proliferated callus on 2N6 media for 30 days"
/dev_stage="proliferated callus on 2N6 media for 30 days"
/lab_host="E.coli DH10B"
/clone lib="Rice callus plasmid cDNA library (NACL)"
/note="Tvector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonucleotides and then used as templates for
RT-PCR."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1M0435A21F Mouse 10kb plasmid UUGCIM library Muclone UUGCIM0435A21 F, genomic survey sequence.
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                                                                                                                                                     High quality sequence stop: 23.
Location/Qualifiers
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                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0435 row: A column: 21
Seq primer: CGTTCTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ610186.1 GI:11732376
                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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primer: SP6; CATACGATTTAGGTGACACTATAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone lib="MPIZ-ADIS-024-developing root"
/note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: Not1;
cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPG-SalI-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
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/mol_type="genomic DNA'
/strain="C57BL/6J"
                                                                      organism="Mus musculus"
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clone="024-026-N02"
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l Similarity
10; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Turkewitz AP
Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 6063
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E., Frankel,J. and Klobutcher,L. EST from Tetrahymena thermophila, strain CU428.1, growing cells Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
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  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA, was prepared from a derivative of physically (gil 4732114 | gb | AF129072.1), a copy-number inducible derivative of plasmid R. The vector was ligated with adaptors complementary to the insert adaptors and nurified the charge desired from the base of the complementary to the insert adaptors.
                                                                             /Strain="CUALO'."
/Strain="CHANO:5911"
/db_xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
preparation can be found in Chilcoat and Turkewitz (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XIIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                              /organism="Tetrahymena thermophila"
/mol_type="mRNA"
/strain="CU428.1"
                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                      0.3%;
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0; Mismatches
                           Score 8.8; I
Pred. No. 12;
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RESULT 12 BM400150

DEFINITION

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SOURCE ORGANISM KEYWORDS

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                                                                              21 bp DNA linear GSS 29
1M0042N20F Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0042N20 F, genomic survey servernament
  Mus
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                                                      AZ321746
AZ321746.1
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI564678 16 bp mRNA linear EST 14 tq78g03.x1 NCI CGAP Ut1 Homo sapiens cDNA clone IWAGE:2214 Similar to TR:Q15214 Q15214 SALIVARY PROLINE-RICH PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.B. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christop
Emmert-Buck, M.D., Ph.D.
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AI564678.1 GI:4523135
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="caxon:9606"
/clone="IMAGE:2214964"
/tissue_type="well-differentiated endometrial
adenocarcinoma, 7 pooled tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="NCI_CGAP_Ut1"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall;
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.75 kb. Life Technologies catalog #:
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                    (house mouse)
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                                                                                                                                                            AZ775540 19 bp DNA linear GSS 16-FEB-200 2M0008H15F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0008H15 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0042 row: N column: 20
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University of Utah Genome
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                               Mus musculus
                                                                                   AZ775540.1
GSS.
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                                                   musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Inducatory Nouse DNA Resource Laboratory Mouse DNA Resources / Occuments / dnares / ). The DNA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF729072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0042N20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sex="Male"
                                                                                                             GI:12902183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 8.8; DB Pred. No. 14; 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 21;
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RESULT 16
CF306647/c
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VERSION
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AUTHORS
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Best Local S
Matches 12
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                                                                                                                                                                                                                                                                                                                                                                                2104 TCCGCTTCCTGCTGGACC 2121
                                                                                                                                                                                                                                                                                                                                          19 recrecrecrecrec 2
                                                                                                                                                                                                                                                                                                                                                                                                                                    1 Similarity
12; Conserv
                                                                                                                                                               CF306647 13 bp mRNA linear EST 15-AUG-2003
HDA1--04-H13.g1 OsHDAC1-overexpressing transgenic rice lambda phage
cDNA library I (HDA1) Oryza sativa cDNA clone HDA1--04-H13, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B.; Hamil Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; 1 (bases 1 to 19)
Oryza sativa
Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0008 row: H column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah
University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                 Oryza sativa
                                                                     CF306647.1
EST.
                                                                                                                    cF306647
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              308,
USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clome_lib="mouse_lokb_plasmid_UUGClm_library" /note="Vector: pWD42nv, Purified genomic_DNA_from_M. musculus_C57BL/6J (male) was obtained_from_the_Jackson_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
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'strain="C57BL/6J"
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Pred. No. 15;
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JOURNAL
                                                                                                                                                                                                MEDLINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches 10;
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Best Local
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                                                                                                                                                                                                                                       JOURNAL
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)

Contact: Nahm B.H.

Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 bp mRNA linear EST 06-DEC-2002
E012766-024-026-N02-SP6 MPIZ-ADIS-024-developing root Beta vulgaris
cDNA clone 024-026-N02 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                              Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                         Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes plant J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                                     and
                                                                                                                                                                                                                                                                                                                    Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Beta vulgaris
Beta vulgaris
            BQ593629.1 GI:26123212
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Oryza sativa"
/mol type="mRNA"
/culfivar="Nackdong"
/db xref="taxon:4530"
/clone="HDA1-04-H13"
/tissue_type="callus"
/dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="E.coli_SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Colone_lib="OSHDAC1-overexpressing transgenic rice lambda phage CDNA library I (HDA1)" (Site 1: EcoRI; Site_2: /notee="Vector: pBluescript SK(+); Site 1: EcoRI; Site_2: XhoI; Callus was treated with ABA(20um) for lhour. cDNA was inserted into lambda Uni-Zap XR vector at 5' end with EcoRI and 3' end with XhoI site. mRNA was derived from rice Histone Deacetylase overexpression line."
    Location/Qualifiers
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Pred. No. 13;
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RESULT 18
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NACL--08-G14.g1 Rice callus plasmid cDNA library (NACL) Oryza
sativa cDNA clone NACL--08-G14, mRNA sequence.
CP332055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 12)
1 (Choses 1, Chong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Sing, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.;
of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CF332055.1 GI:33812331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: bhnahm@ggbio.com, bhnahmannahm@ggbio.com, bhnahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannahmannah
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone lib="MPIZ-ADIS-024-developing root"
/clone lib="MPIZ-ADIS-024-developing root"
/note="Vector: pcMVSPORT6; Site 1: SalI; Site 2: NotI;
/note="Vector: pcMVSPORT6; Site 1: SalI; Site 2: NotI;
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/note="Vector: pcMVSPORT6; SalI-NotI; primer sites and bischulz@kws.de; cloning sites SalI-NotI, primer sites and bischulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   / moi_type="mRNA"
/cultivar-"-
                                                                       /cultivar="Nackdong"
/db xref="Nackdong"
/db xref="Nackdong"
/db xref="Nackdong"
/clone="NaCL--08-014"
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/clone_lib="Rice_callus plasmid cDNA library (NACL)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Beta vulgaris"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:161934"
/clone="024-026-N02"
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/lab_host="EMDH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       db_xref="GABI:193221"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Oryza sativa"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Korea
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Pred. No. 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bhnahm@bio.myongji.ac.kr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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Length 12;
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Query Match Best Local Similarity

0.3%;

Score 7.8; DB 1; Pred. No. 13;

Matches	9;	Conserva	tive		9; Conservative 0; Mismatches	2;	2; Indels 0; Gaps 0;	0;	Gaps	0;
	3 	1603 CCCCGTGCTTC 1613	1613							
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search completed: April 7, 2004, 16:19:57 Tob time : 1 secs	secs	April '	7, 2004,	16	:19:57					